

● PRINTER RUSH ●
(PTO ASSISTANCE)

Application :	<u>10/002,750</u>	Examiner :	<u>Burkhart</u>	GAU :	<u>1636</u>
From:	<u>J. Robbins</u>	Location:	<u>IDC</u> FMF FDC	Date:	<u>9-7-05</u>
Tracking #:			<u>06094328</u>	Week Date: <u>4-11-05</u>	

DOC CODE	DOC DATE	MISCELLANEOUS
<input type="checkbox"/> 1449	_____	<input type="checkbox"/> Continuing Data
<input type="checkbox"/> IDS	_____	<input type="checkbox"/> Foreign Priority
<input type="checkbox"/> CLM	_____	<input type="checkbox"/> Document Legibility
<input type="checkbox"/> IIFW	_____	<input type="checkbox"/> Fees
<input type="checkbox"/> SRFW	_____	<input type="checkbox"/> Other
<input checked="" type="checkbox"/> DRW	<u>6-21-02</u>	
<input type="checkbox"/> OATH	_____	
<input type="checkbox"/> 312	_____	
<input type="checkbox"/> SPEC	_____	

[RUSH] MESSAGE: ATTN: Chief drafts person

Figs 1-4, 6, 37A and 37B contain
a line through drawings.

Thank You
[Signature]

[XRUSH] RESPONSE: _____

DRAWINGS CORRECTED

9-13-2005

INITIALS [Signature]

NOTE: This form will be included as part of the official USPTO record, with the Response document coded as XRUSH.
REV 10/04

COMPLEMENTING CELL LINES

Inventor: Vogels et al
 Serial No.: 10/002,750
 Docket No.: 2183-5148US

Figure 1:
 % of human sera with neutralising capacity for human adenovirus (n=100)

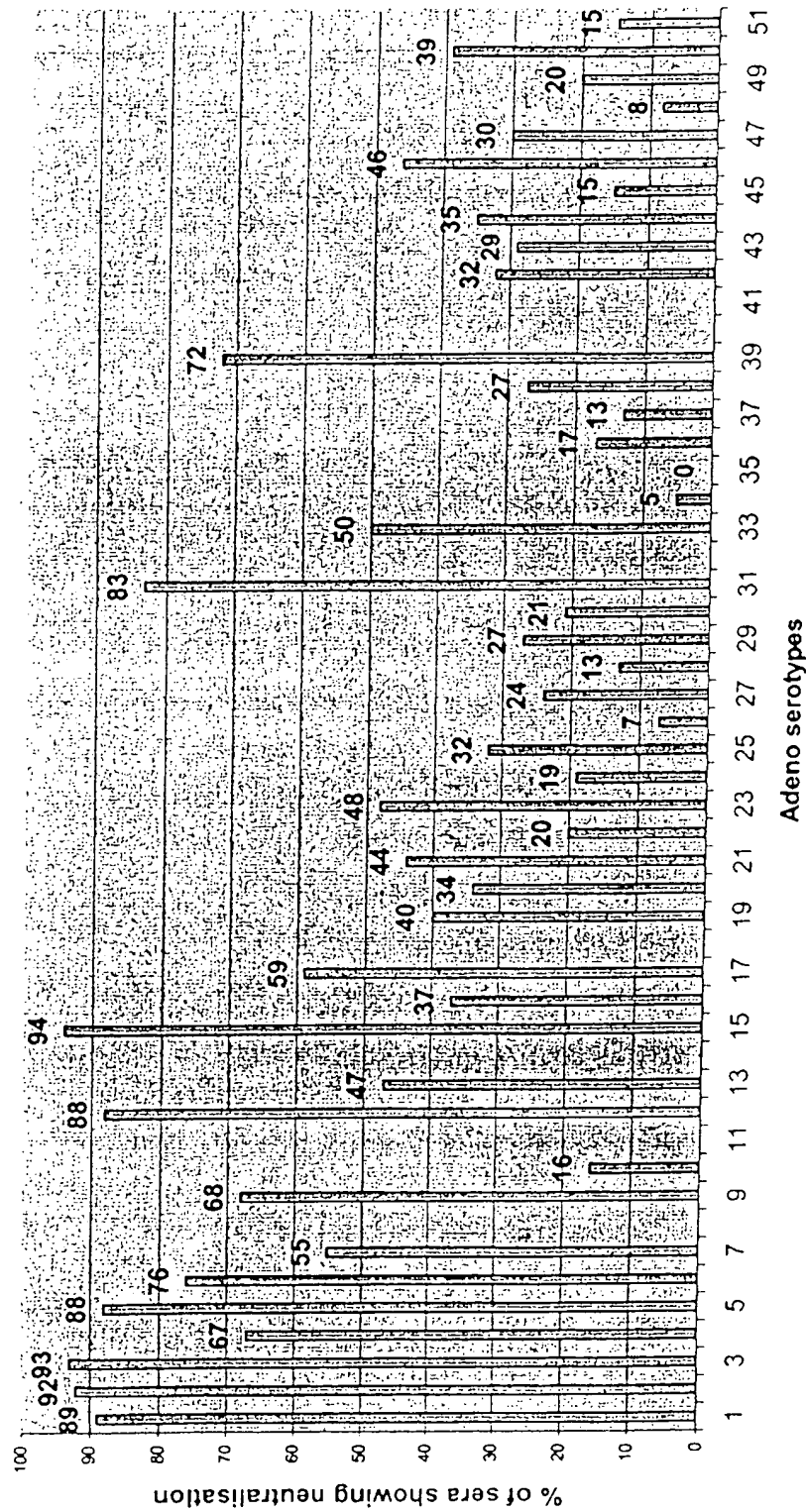


Figure 2

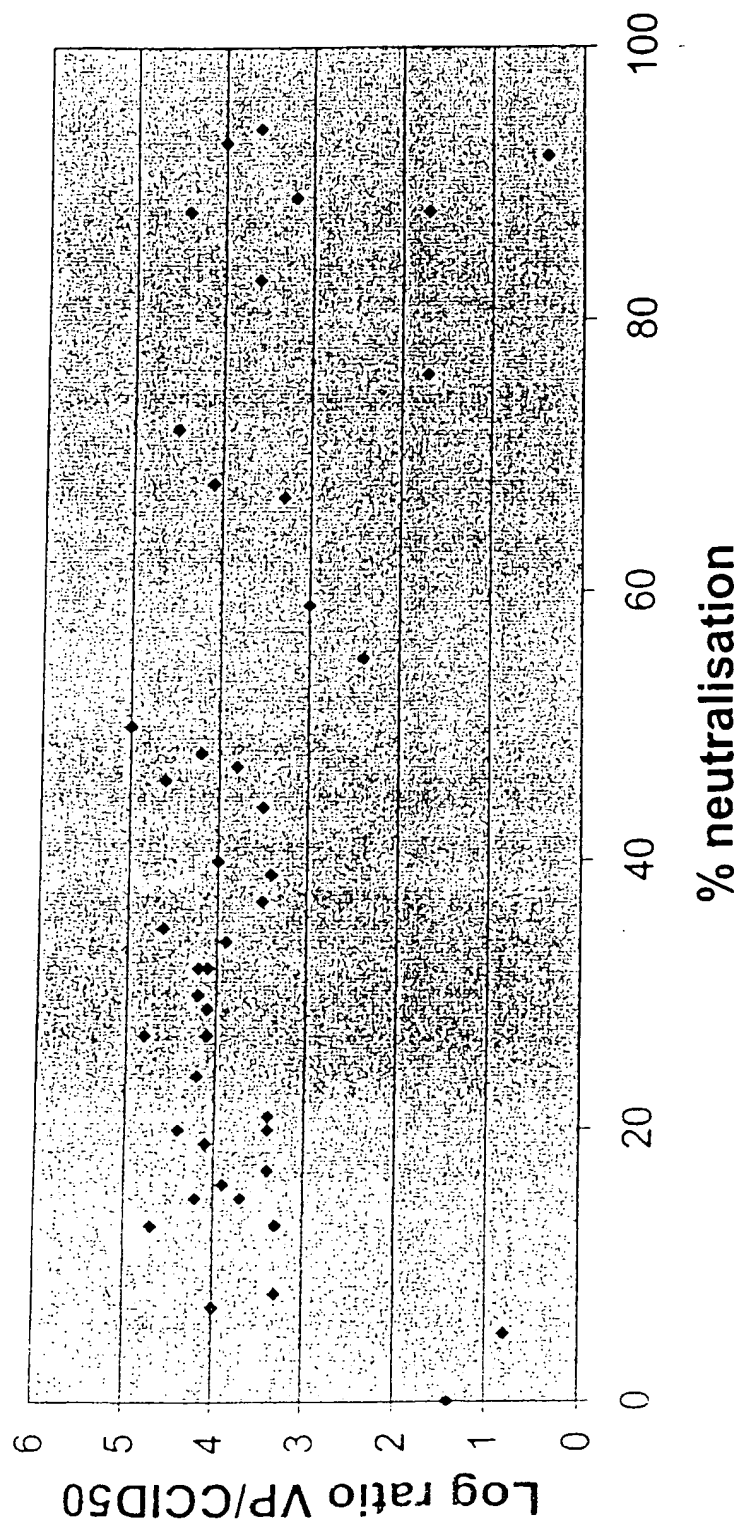


Figure 3
 Neutralisation in human sera

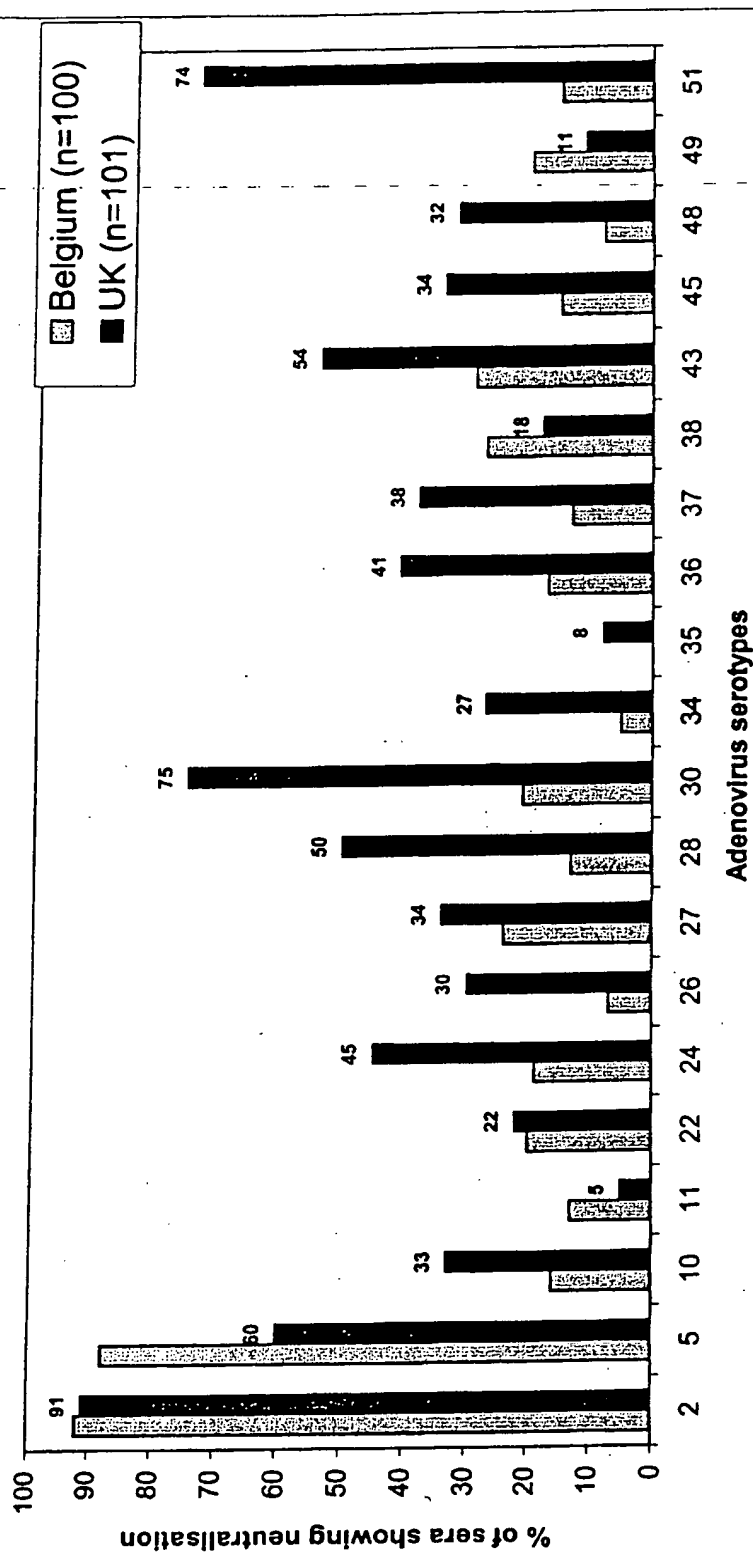


Figure 4
Neutralisation in human sera from different geographic locations

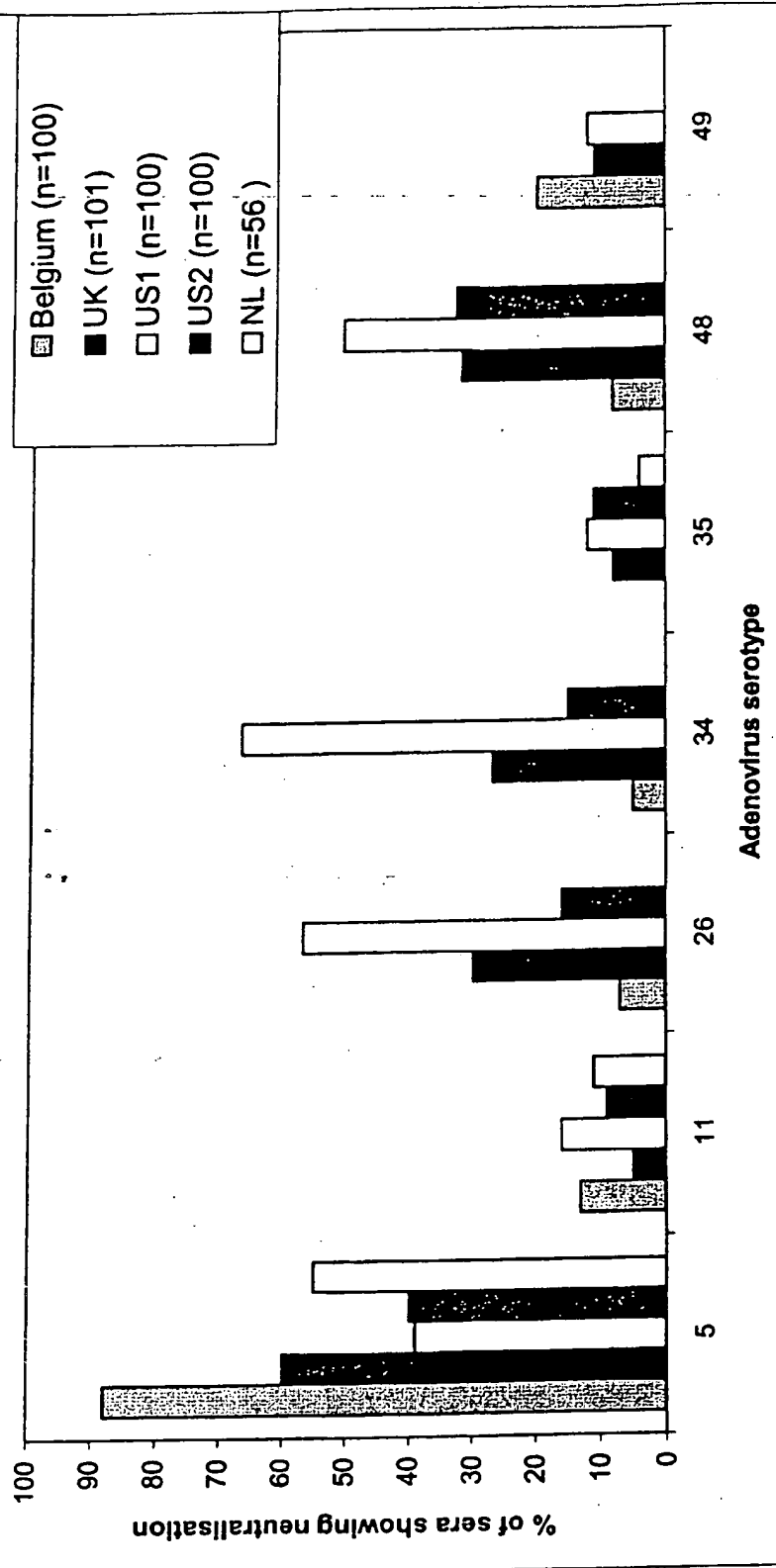


Figure 5

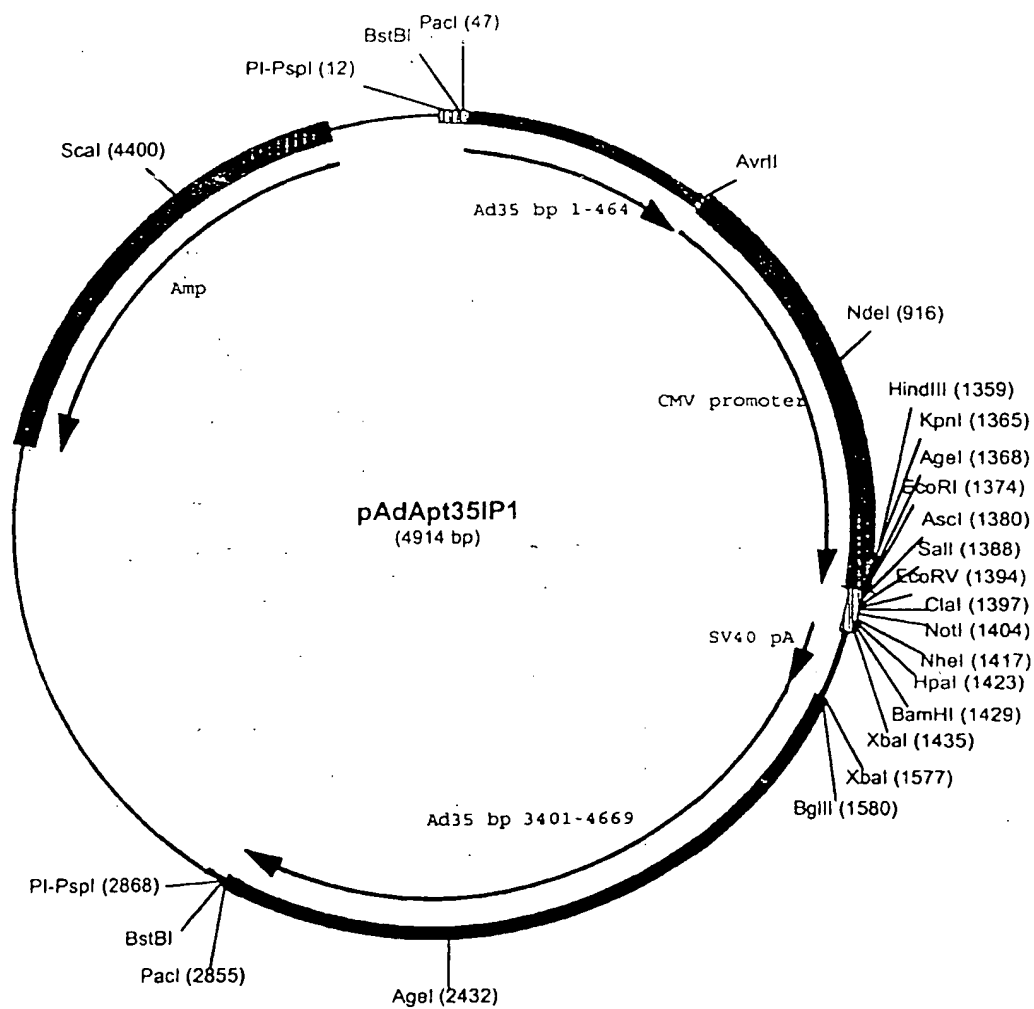


Figure 6

Construction of cosmid vector pWE.Ad35.pIX-rITR

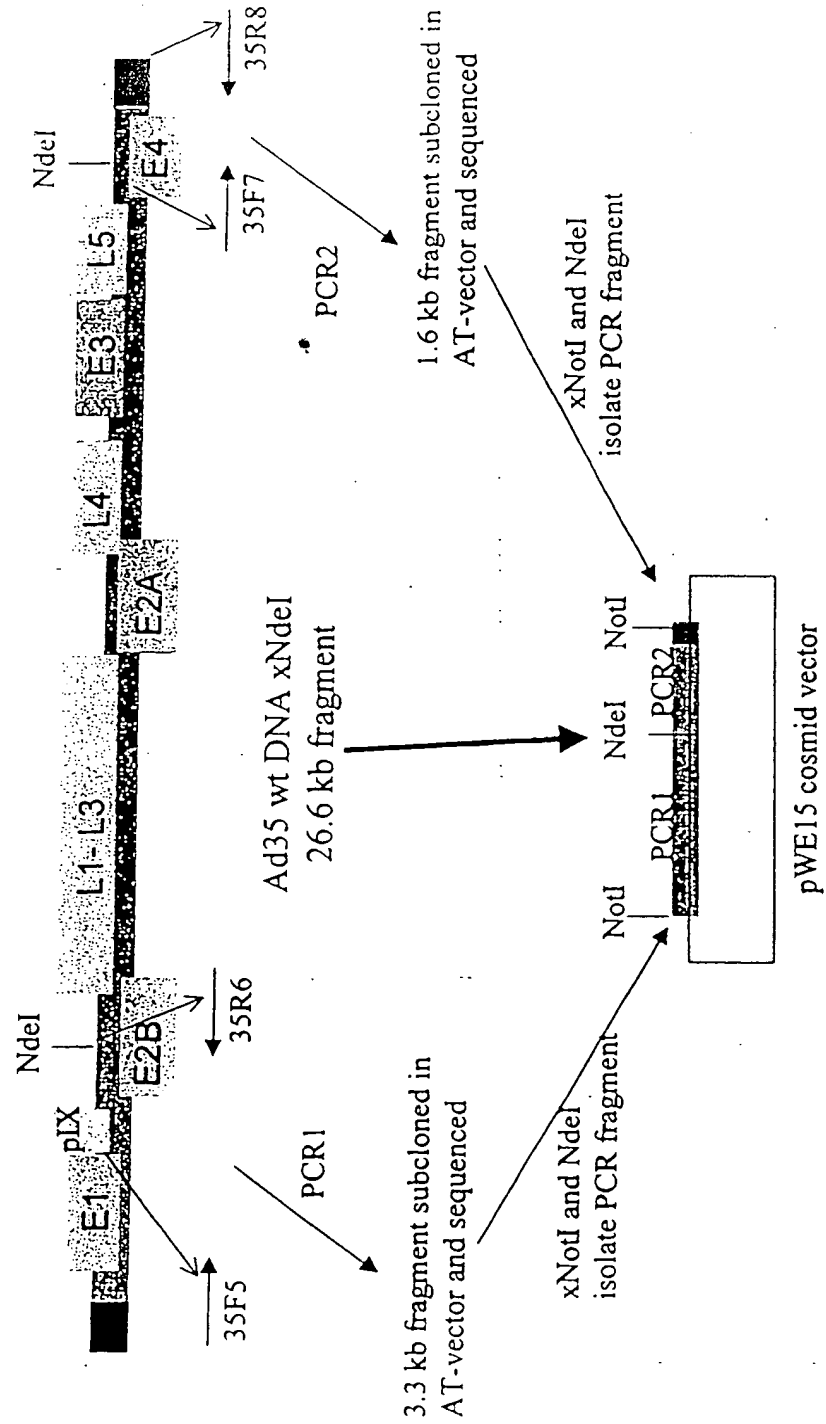


Figure 7

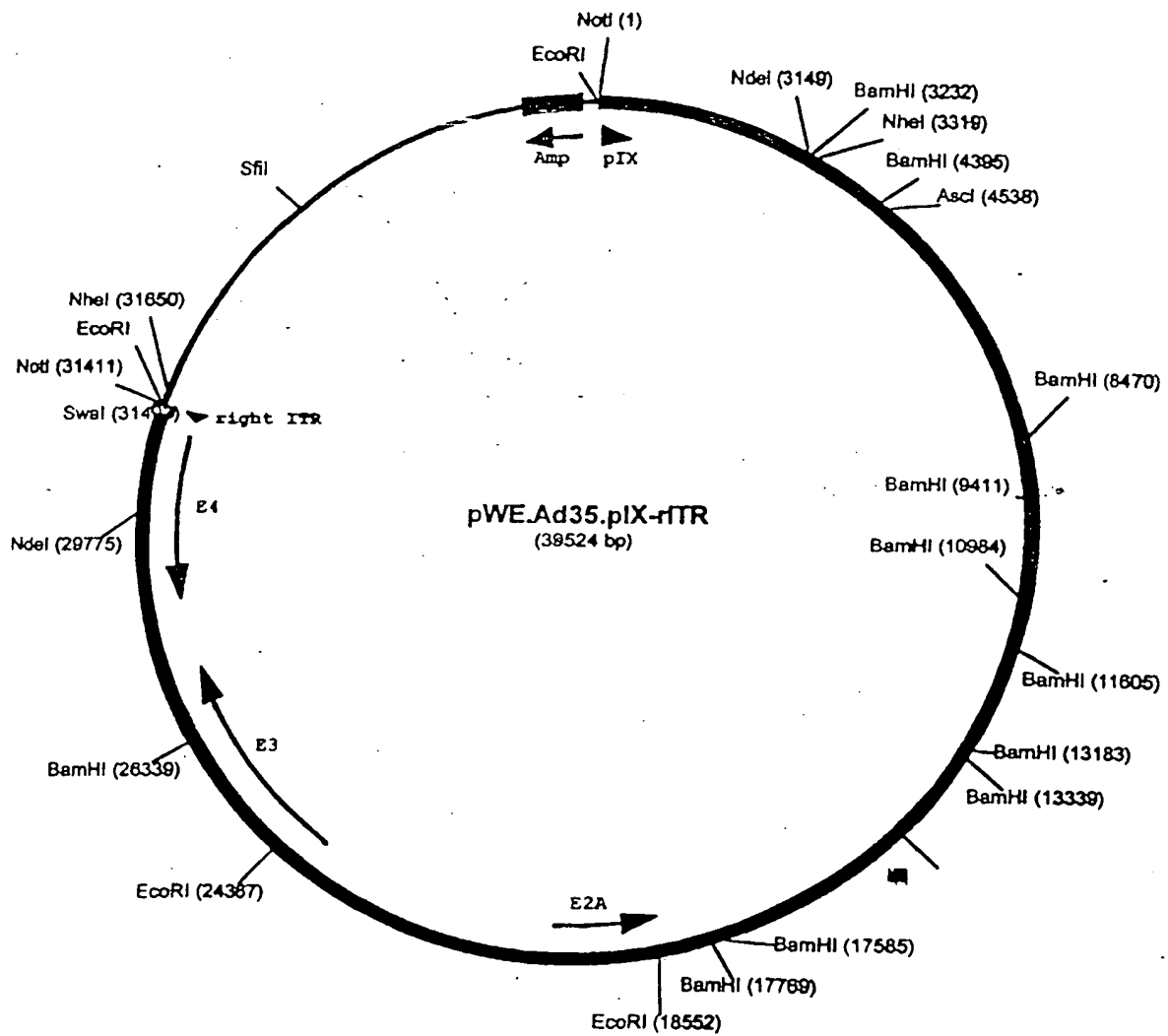


Figure 8

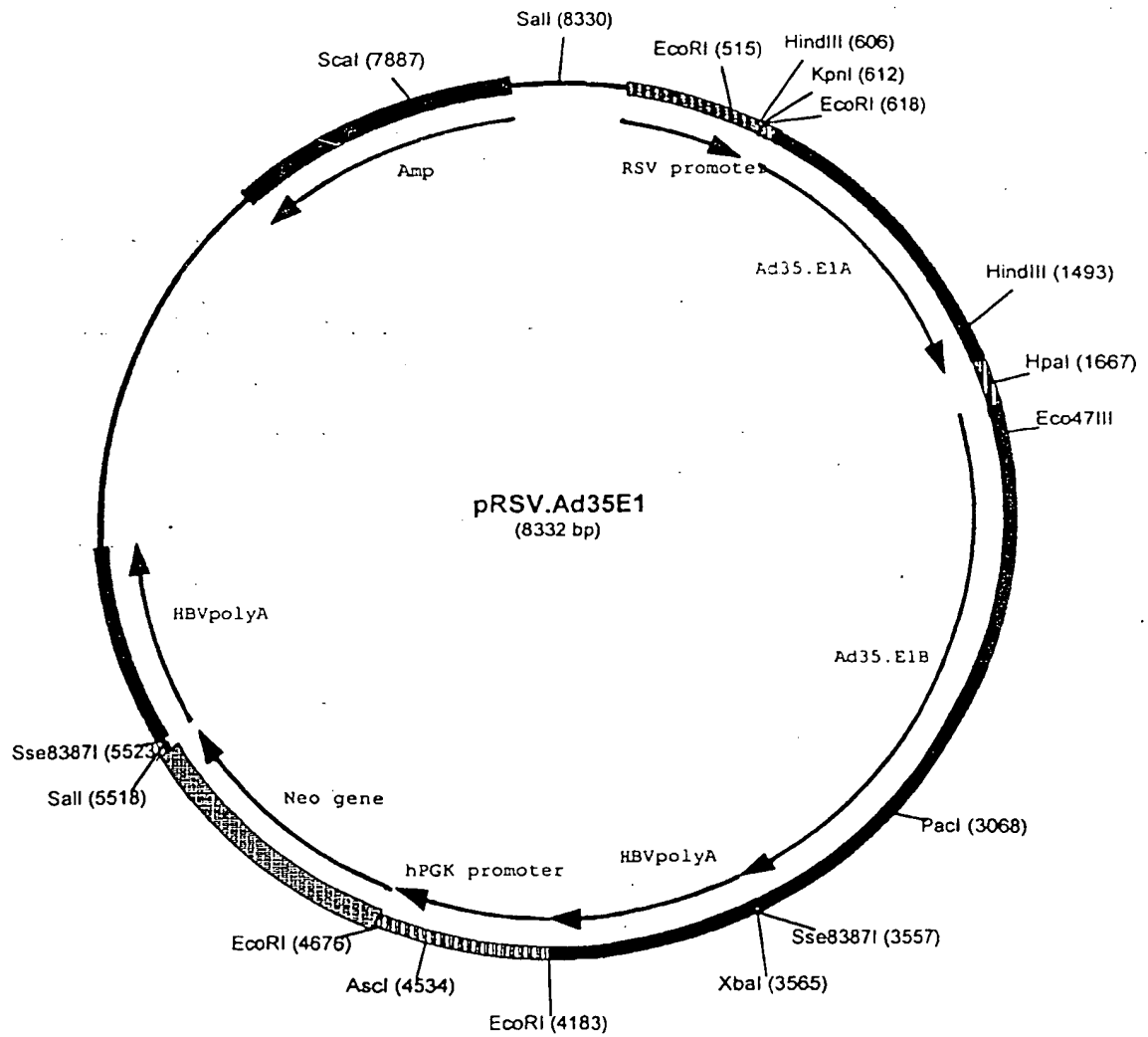


Figure 9

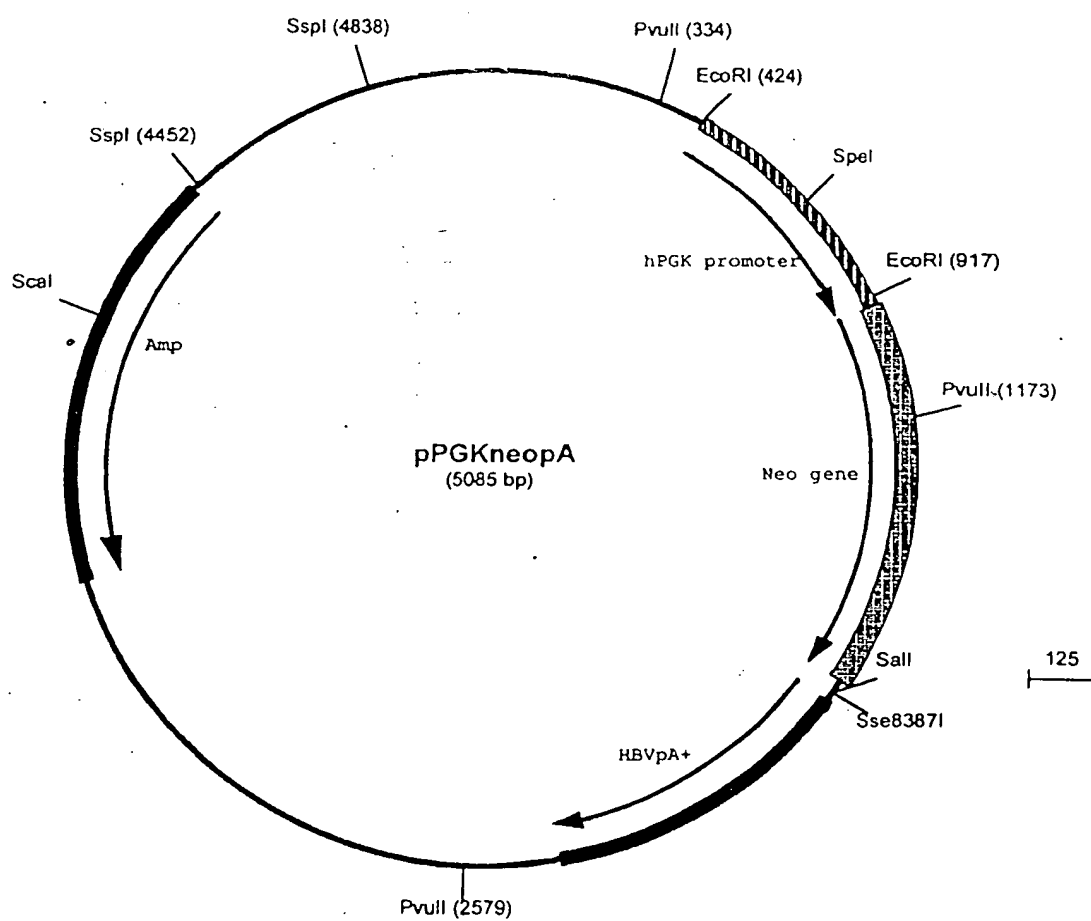


Figure 10

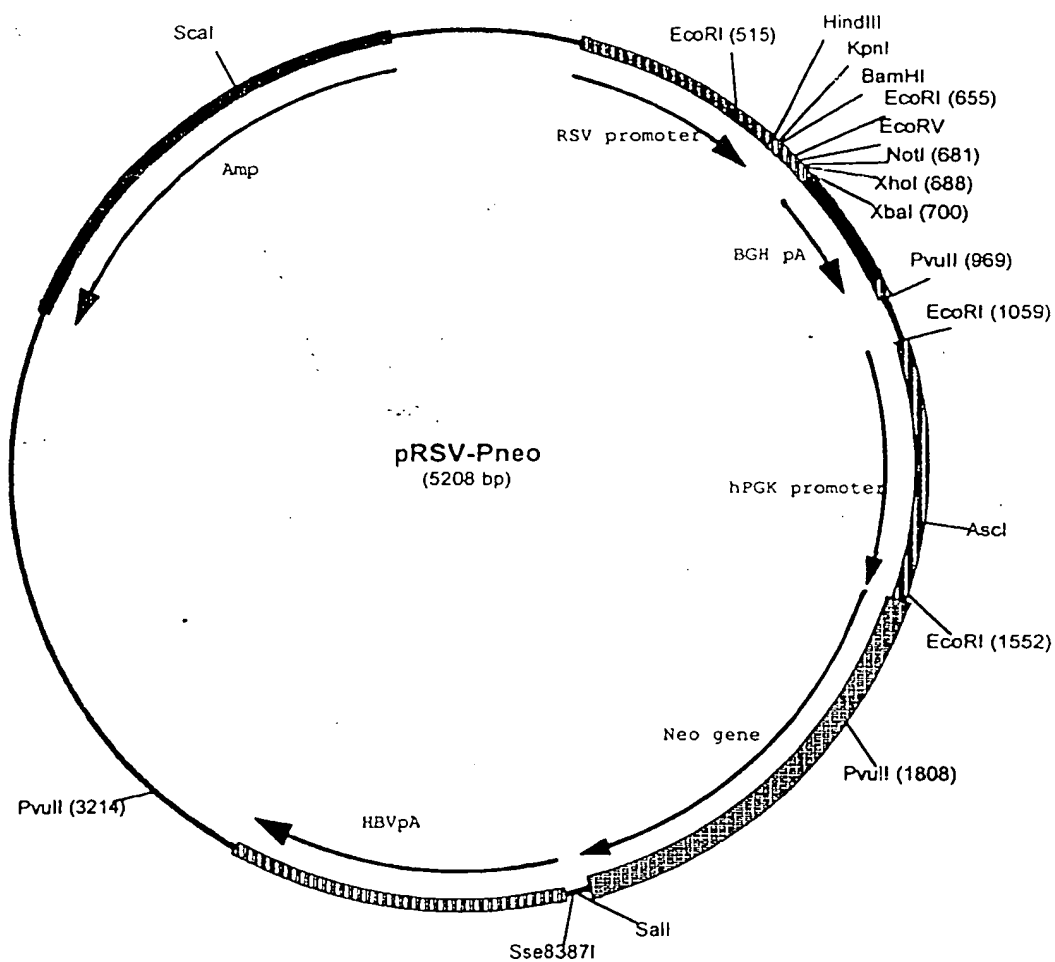


Figure 11

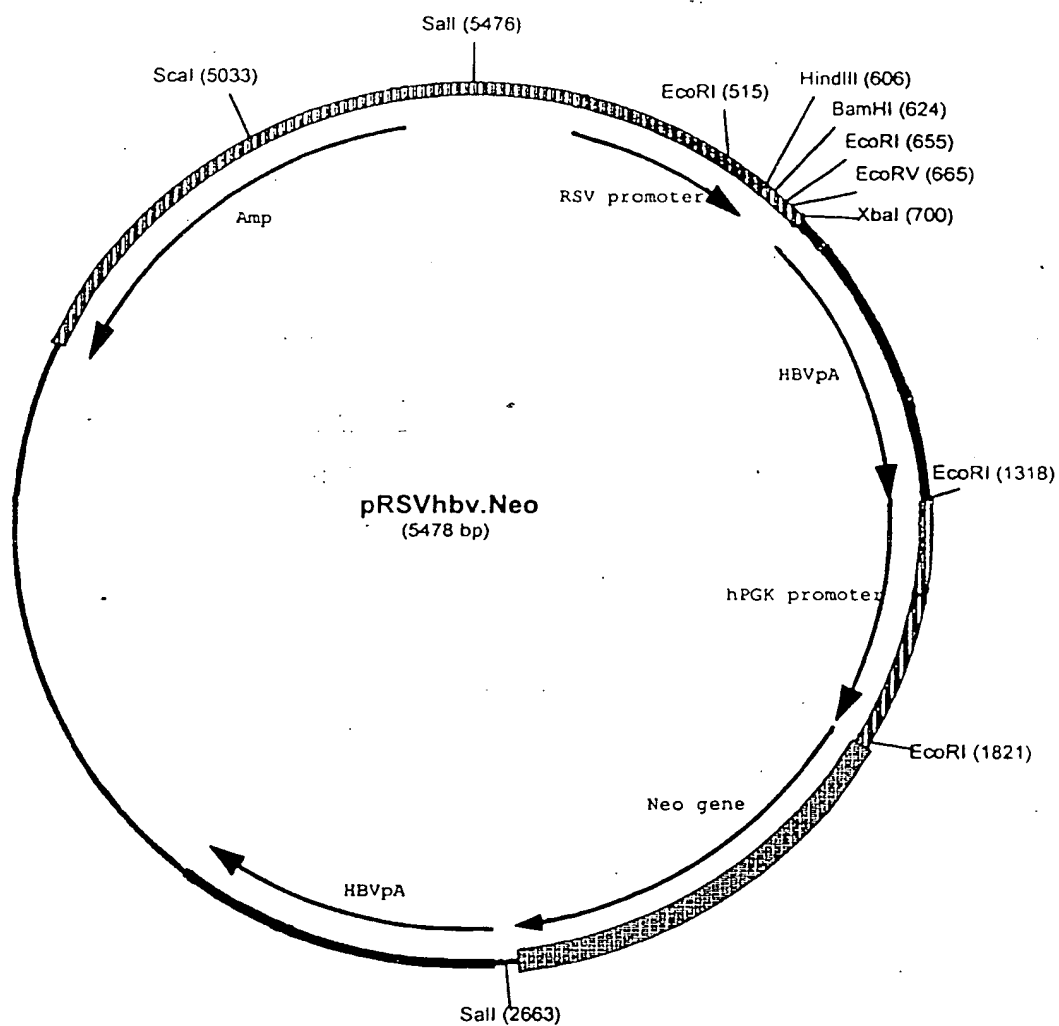


Figure 12

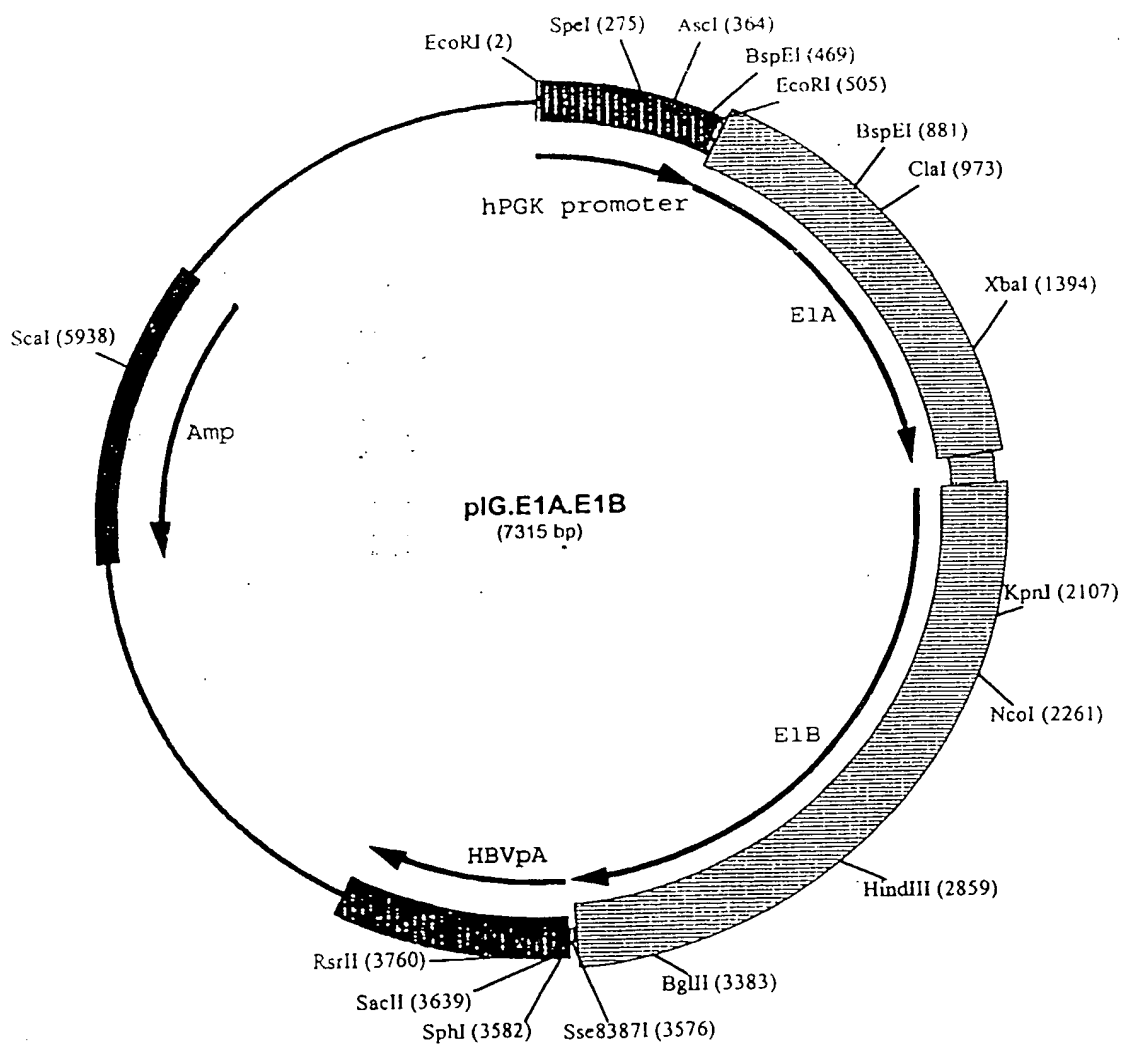


Figure 13

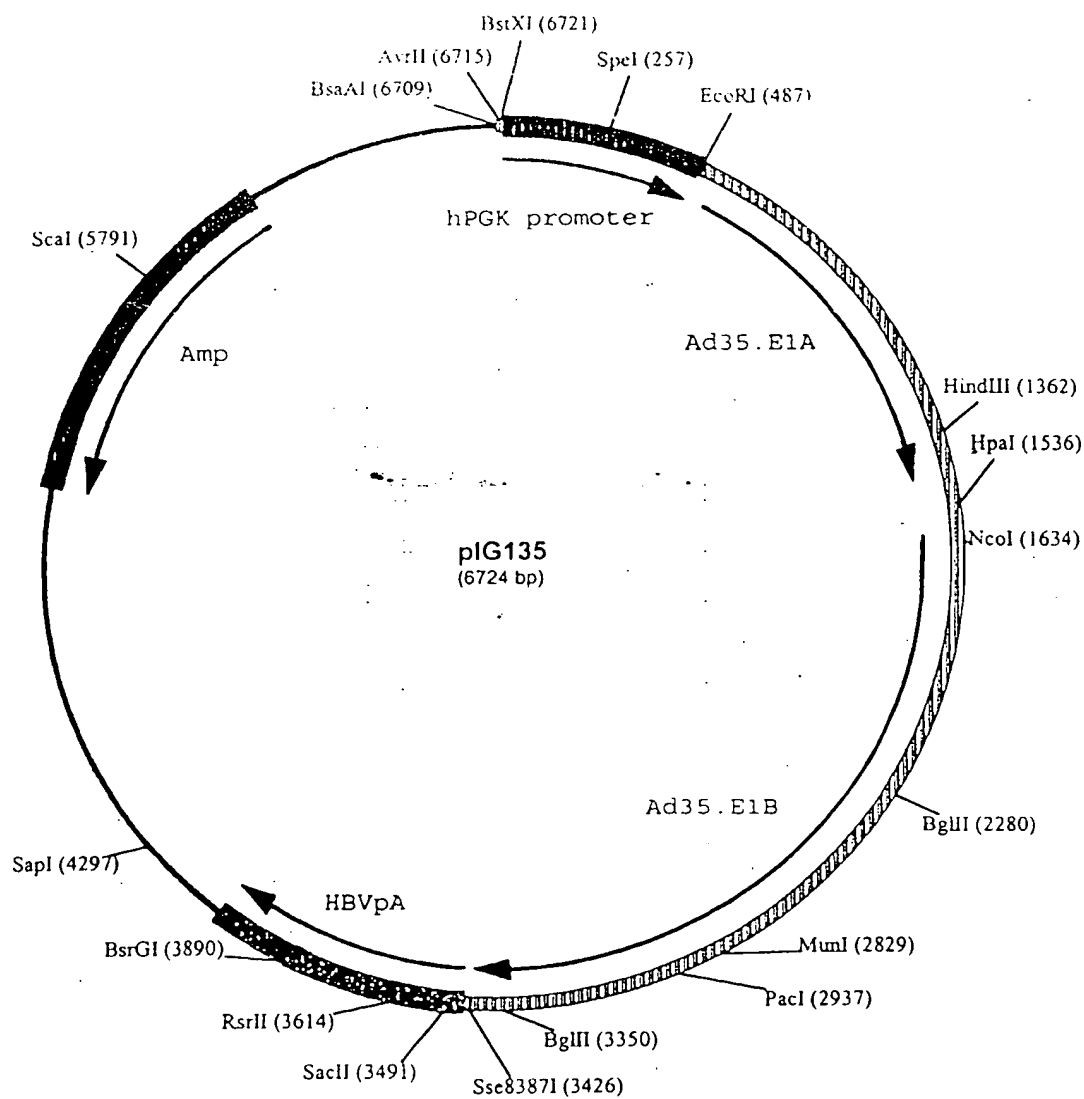


Figure 14

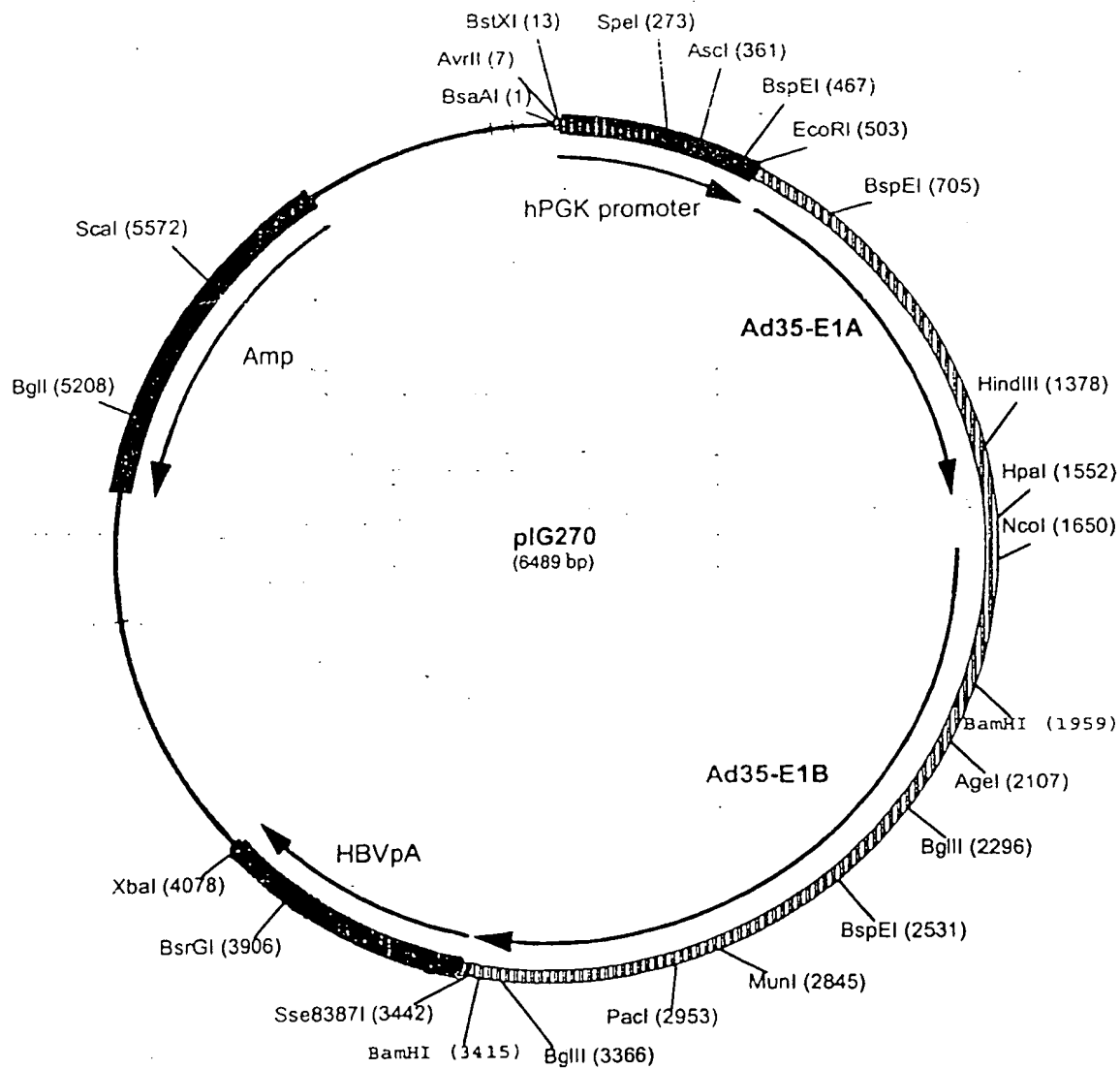


Figure 15

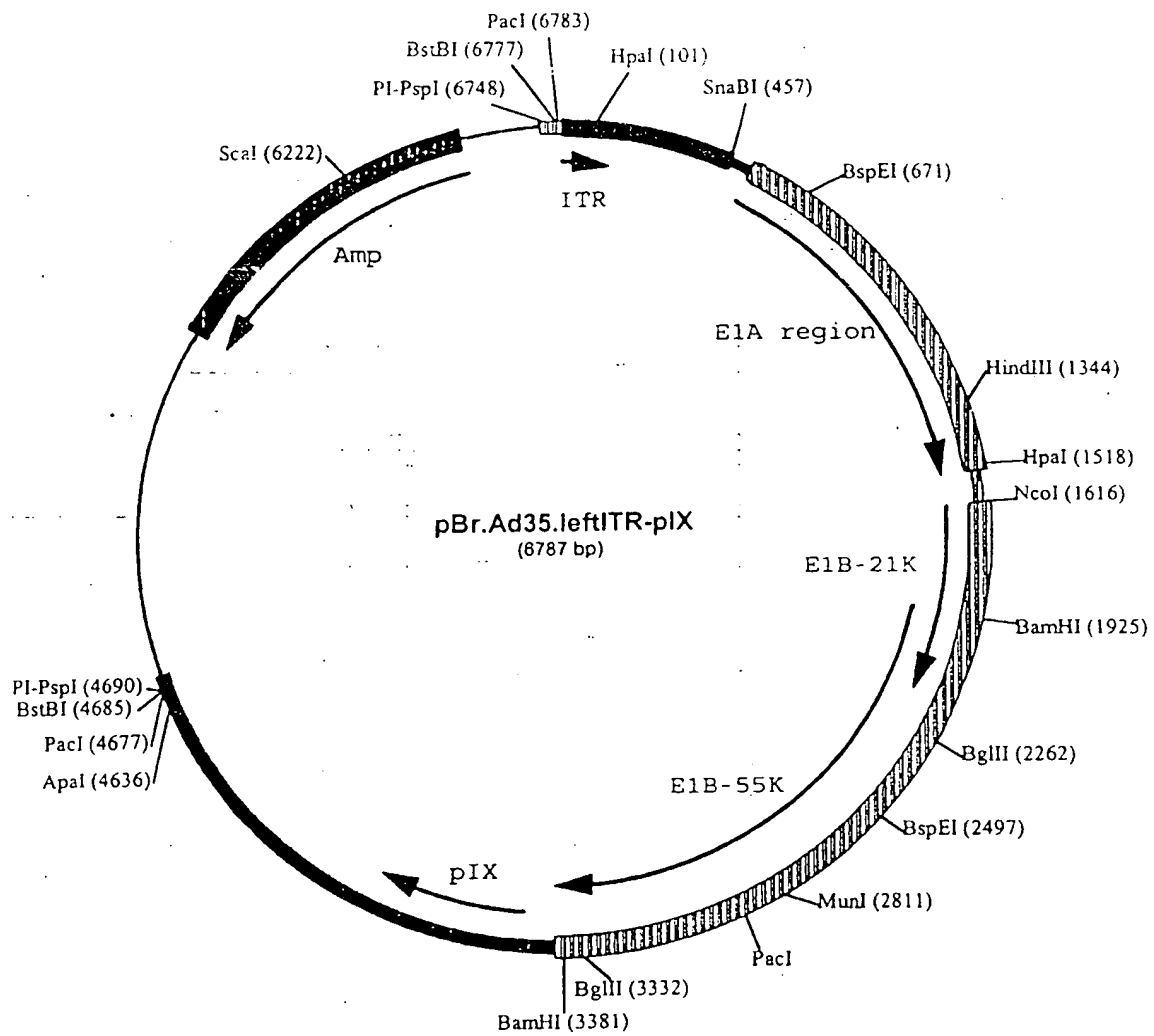


Figure 16

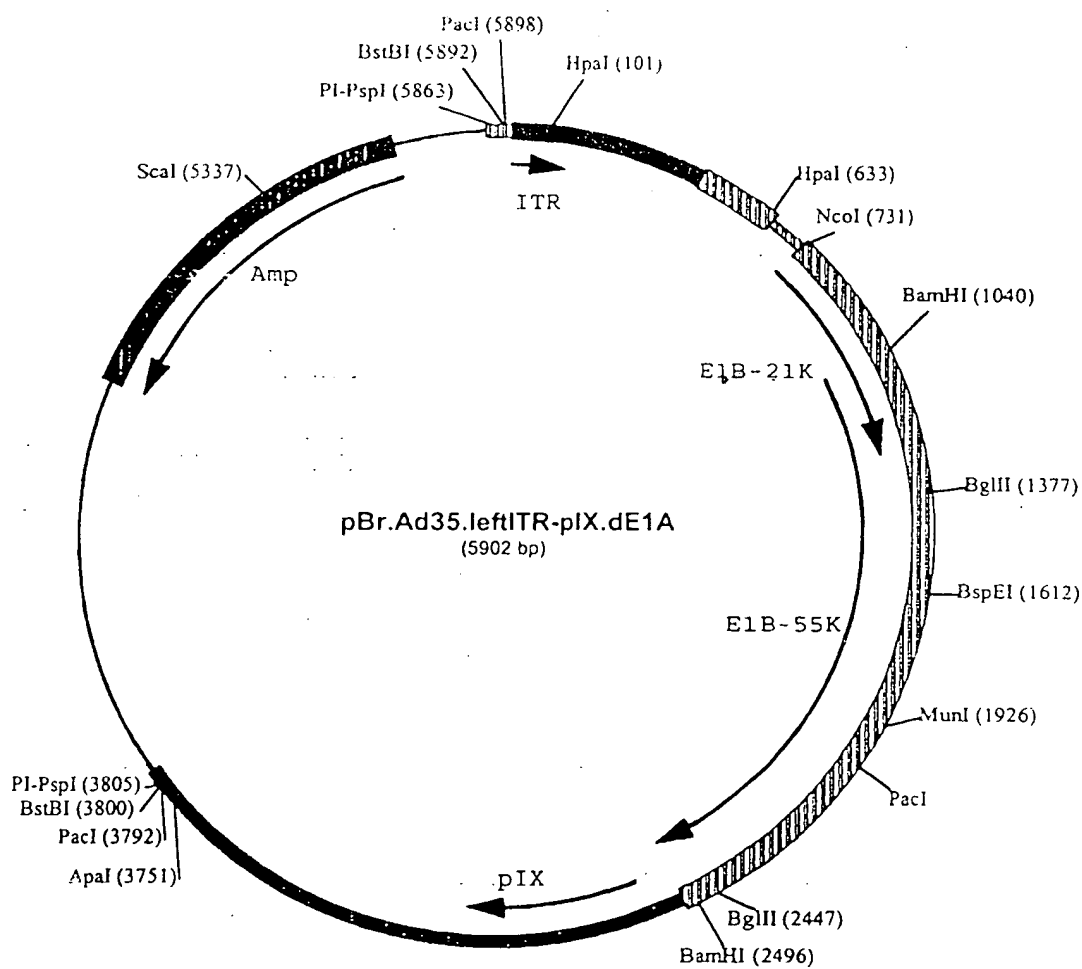


Figure 17

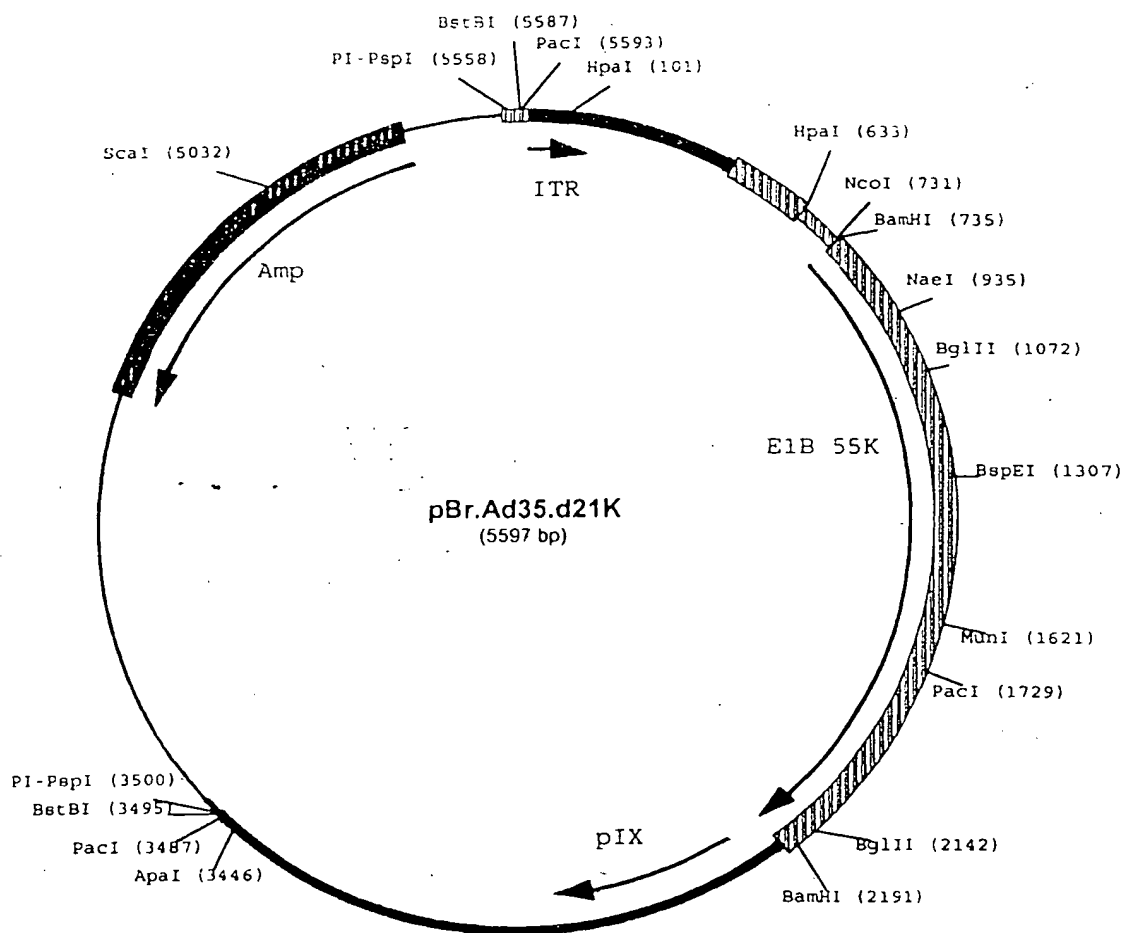


Figure 18

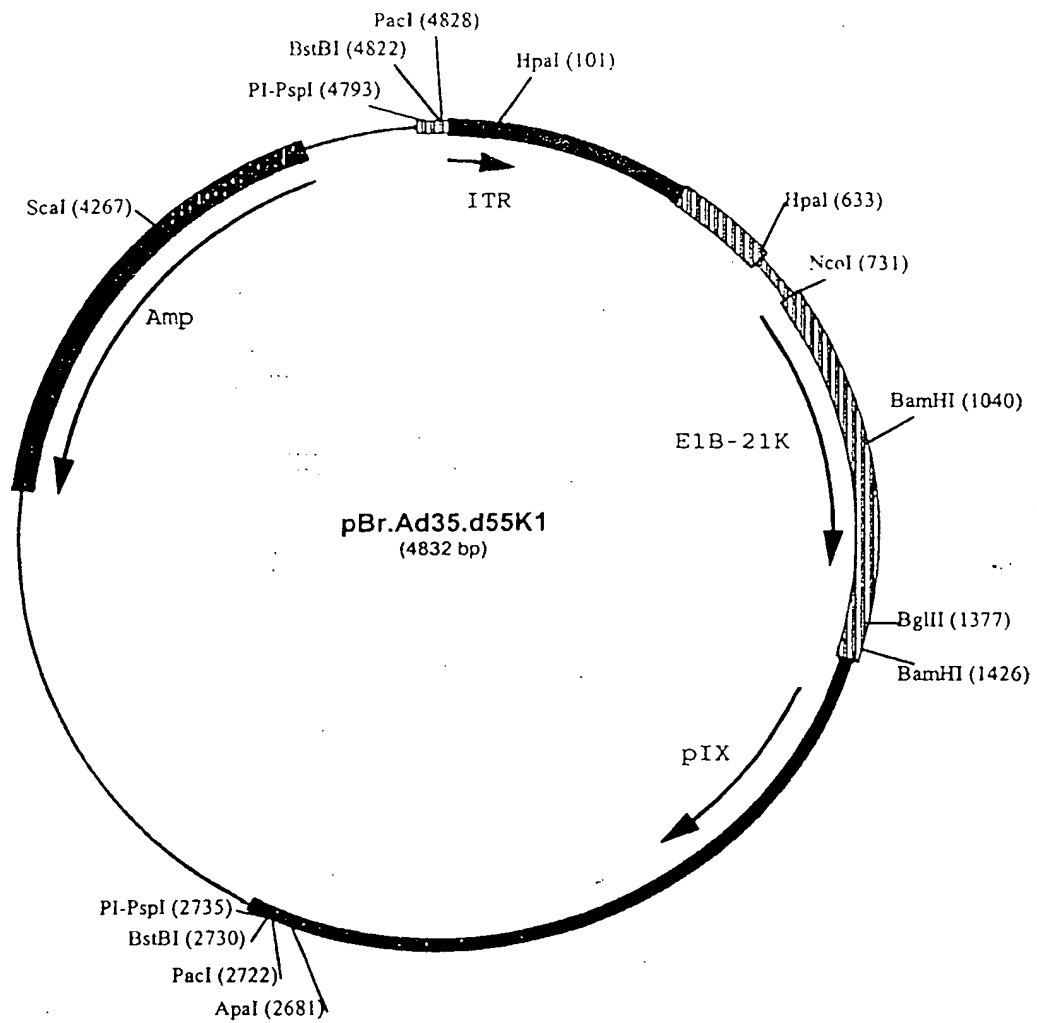


Figure 19

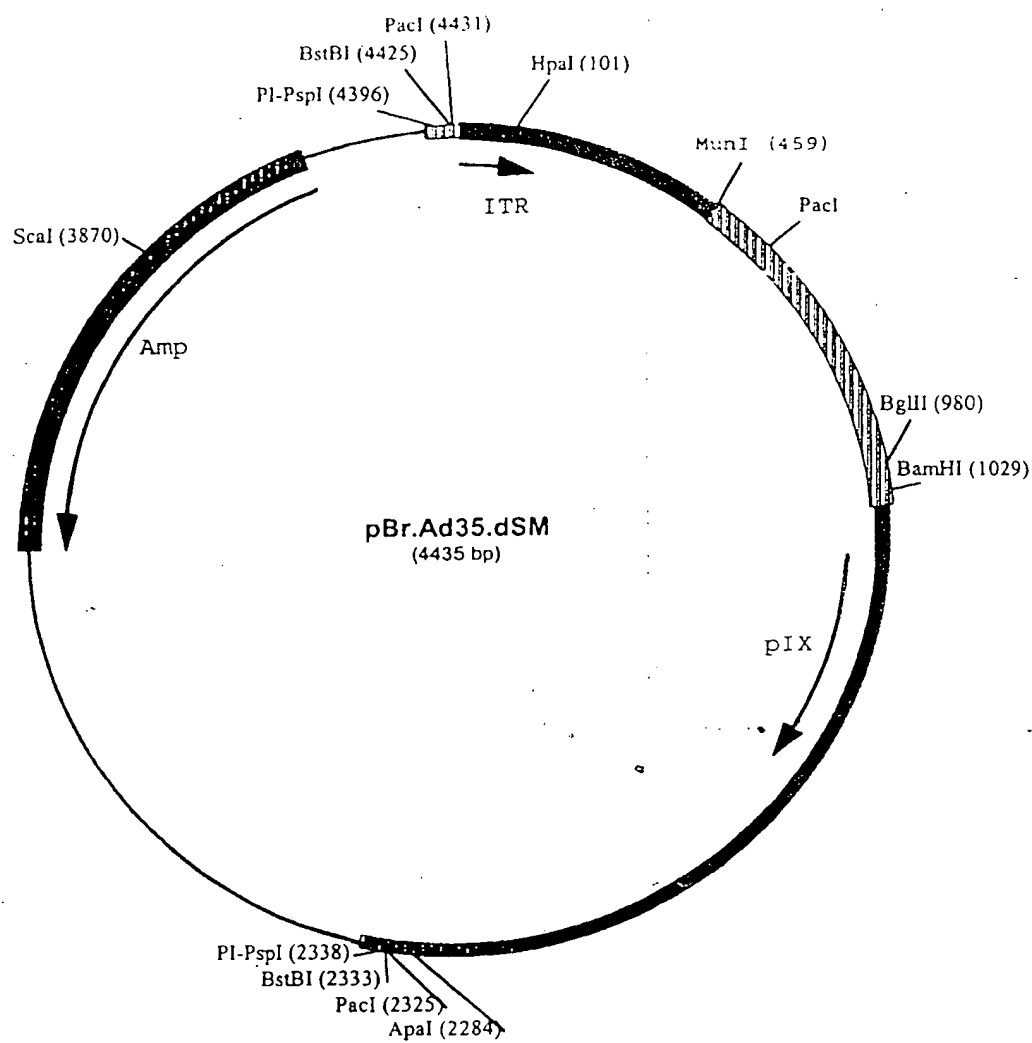


Figure 20

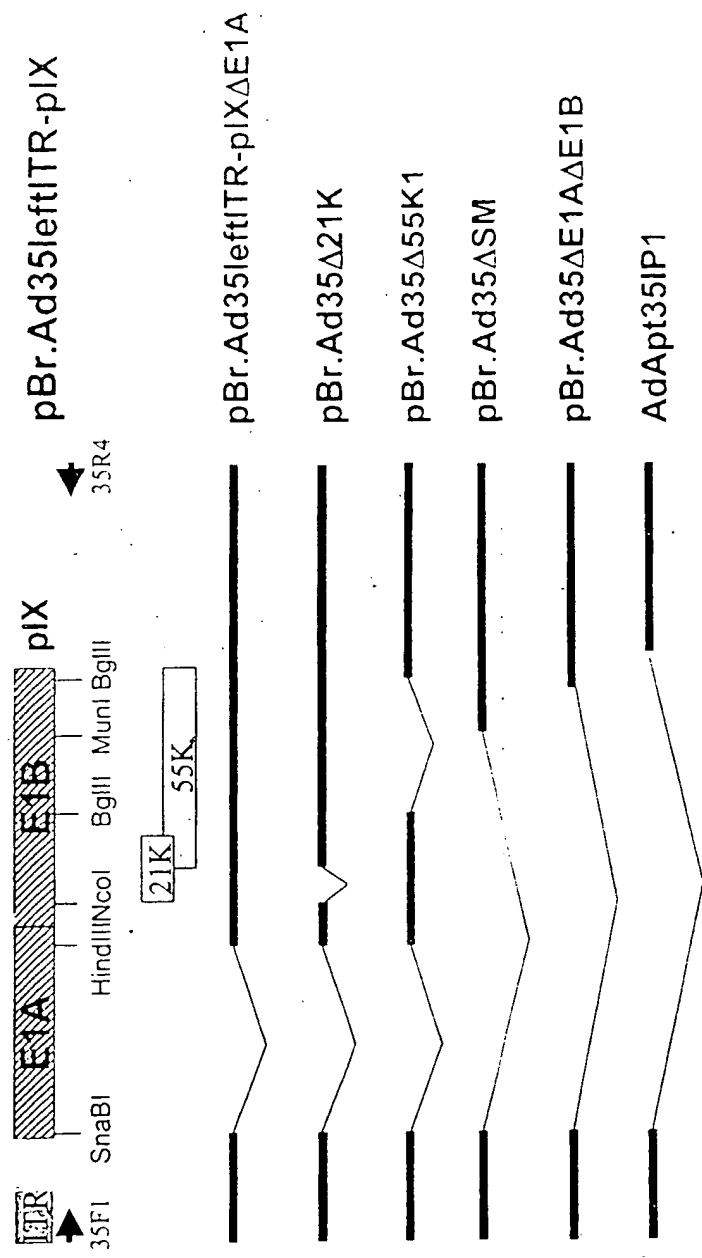


Figure 21

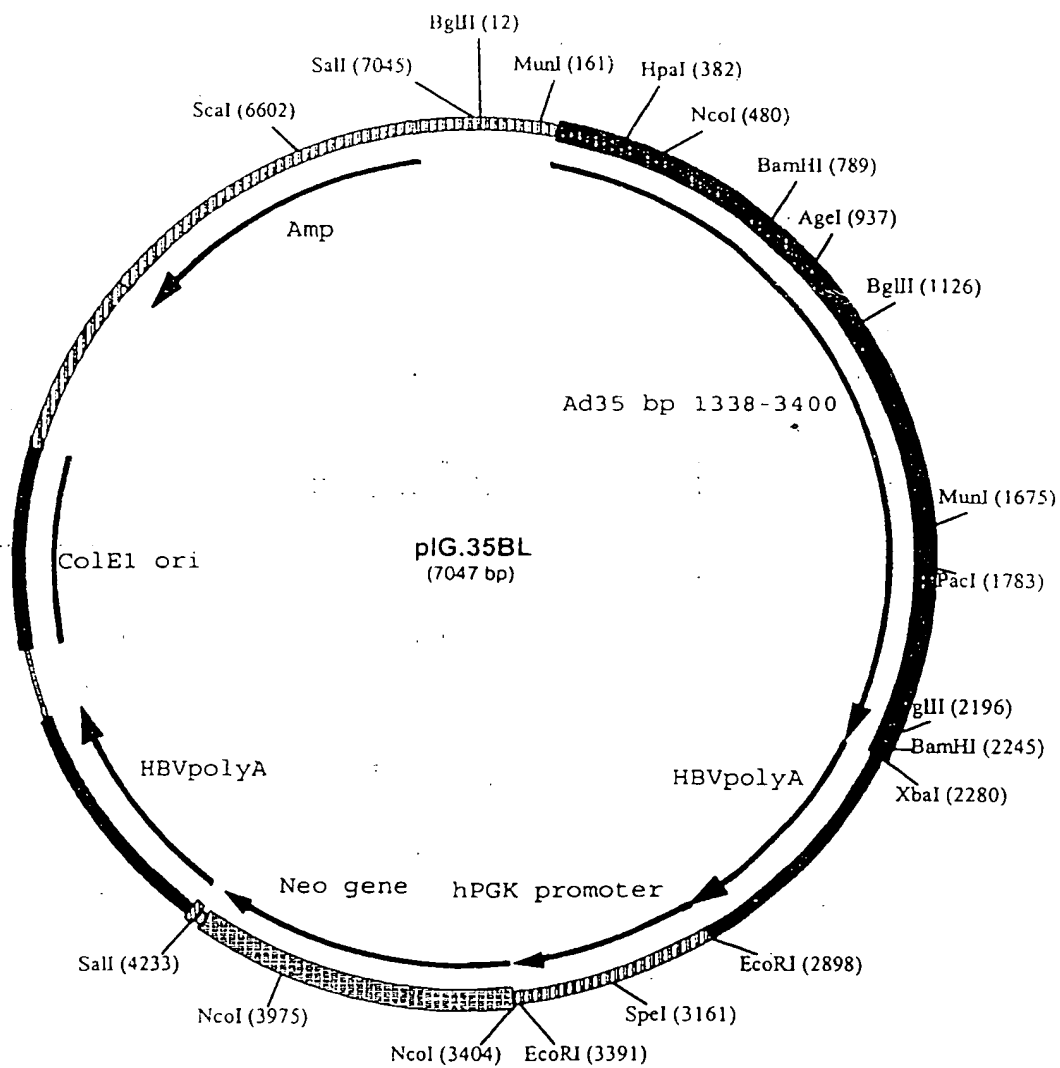


Figure 22

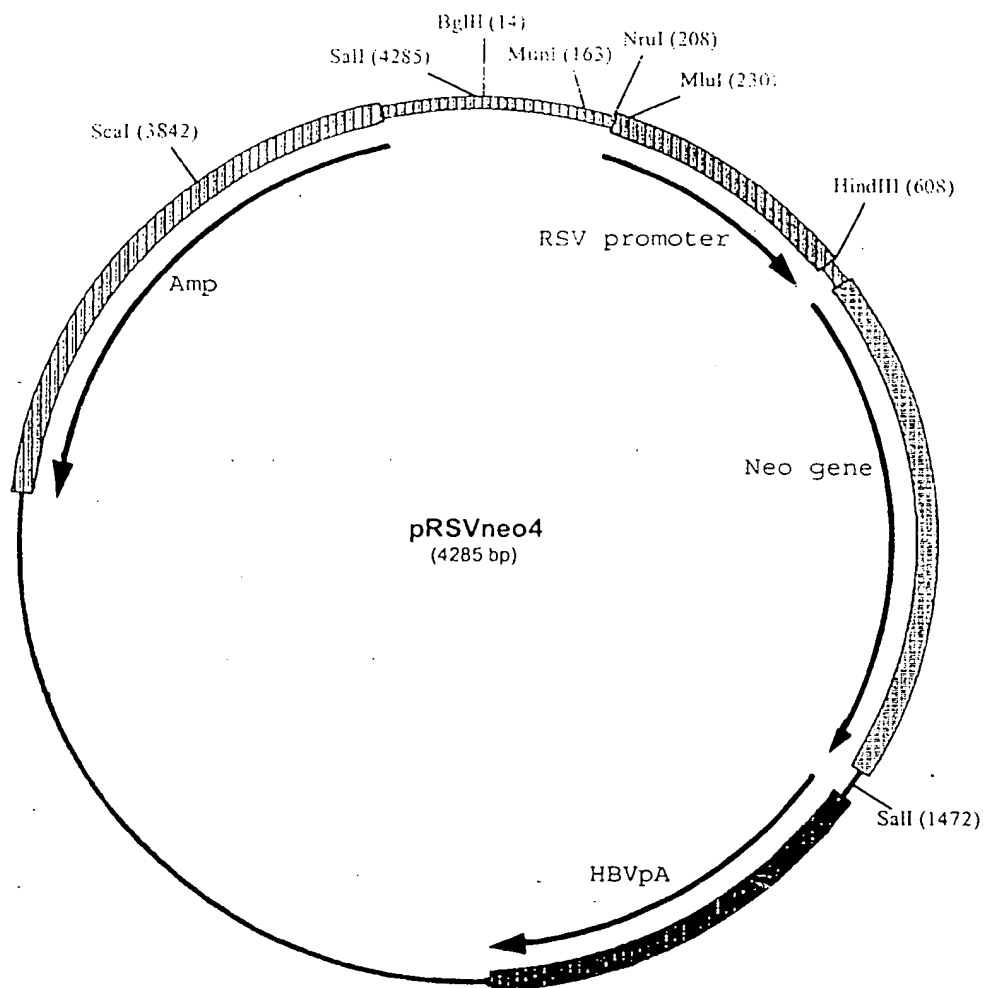


Figure 23

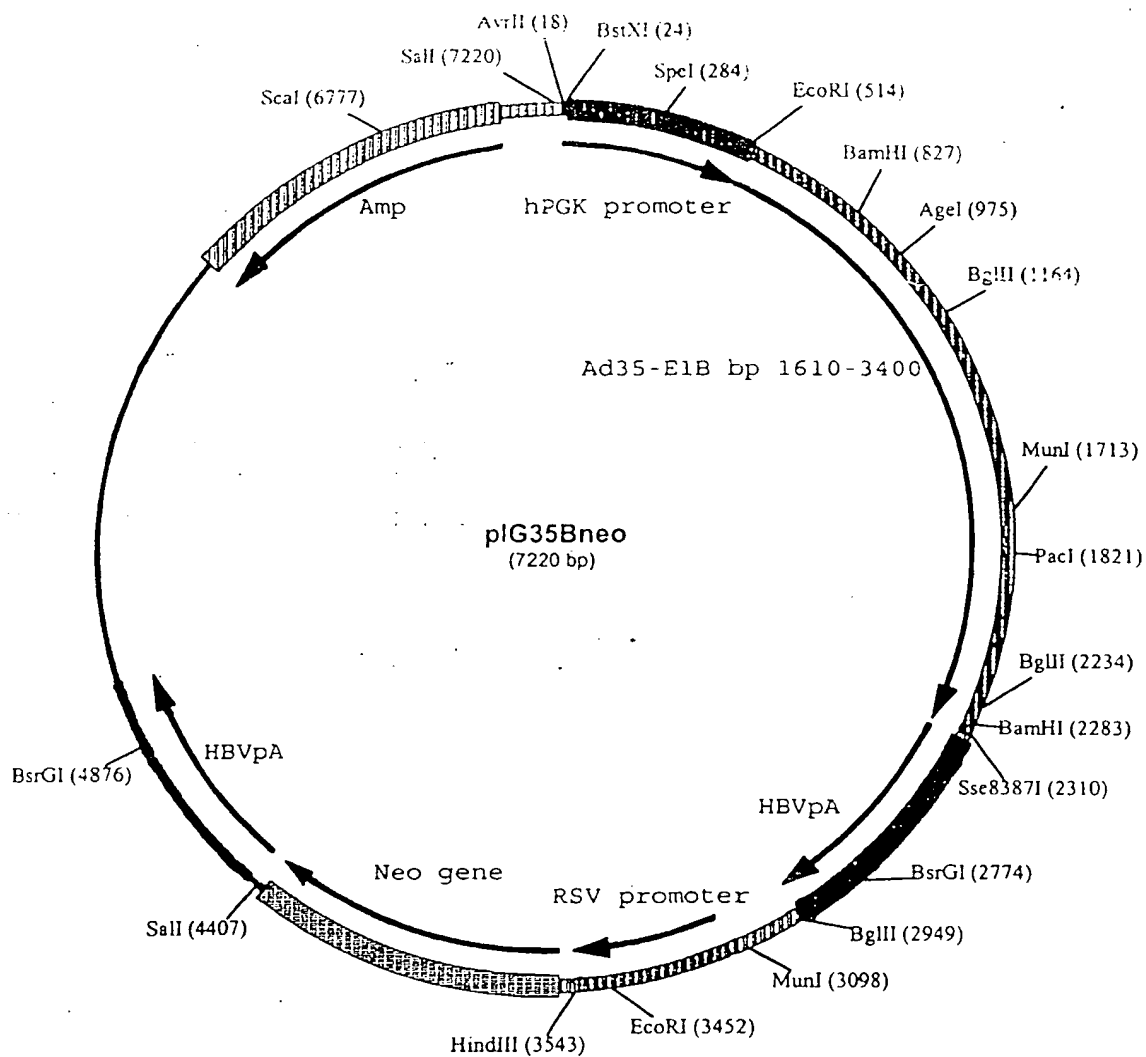


Figure 24

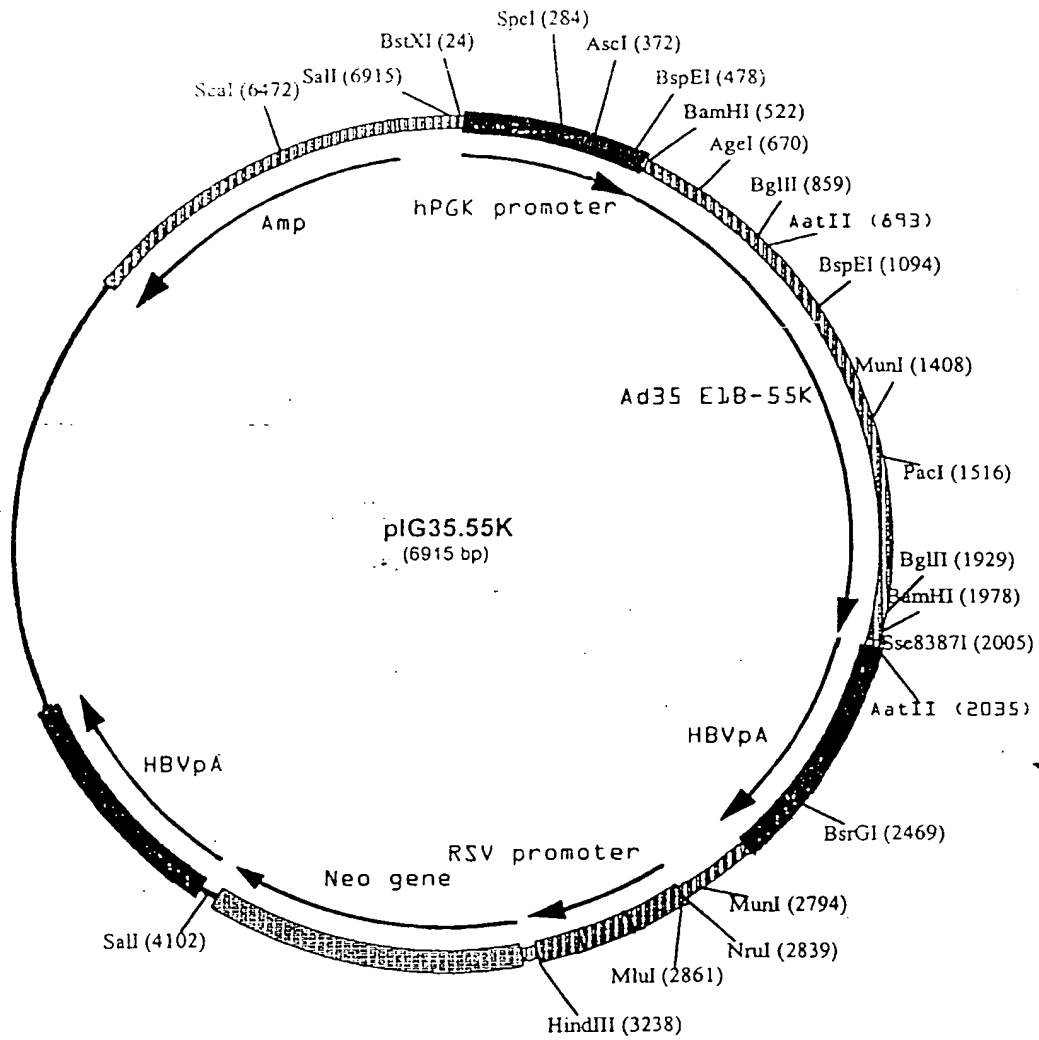


Figure 25

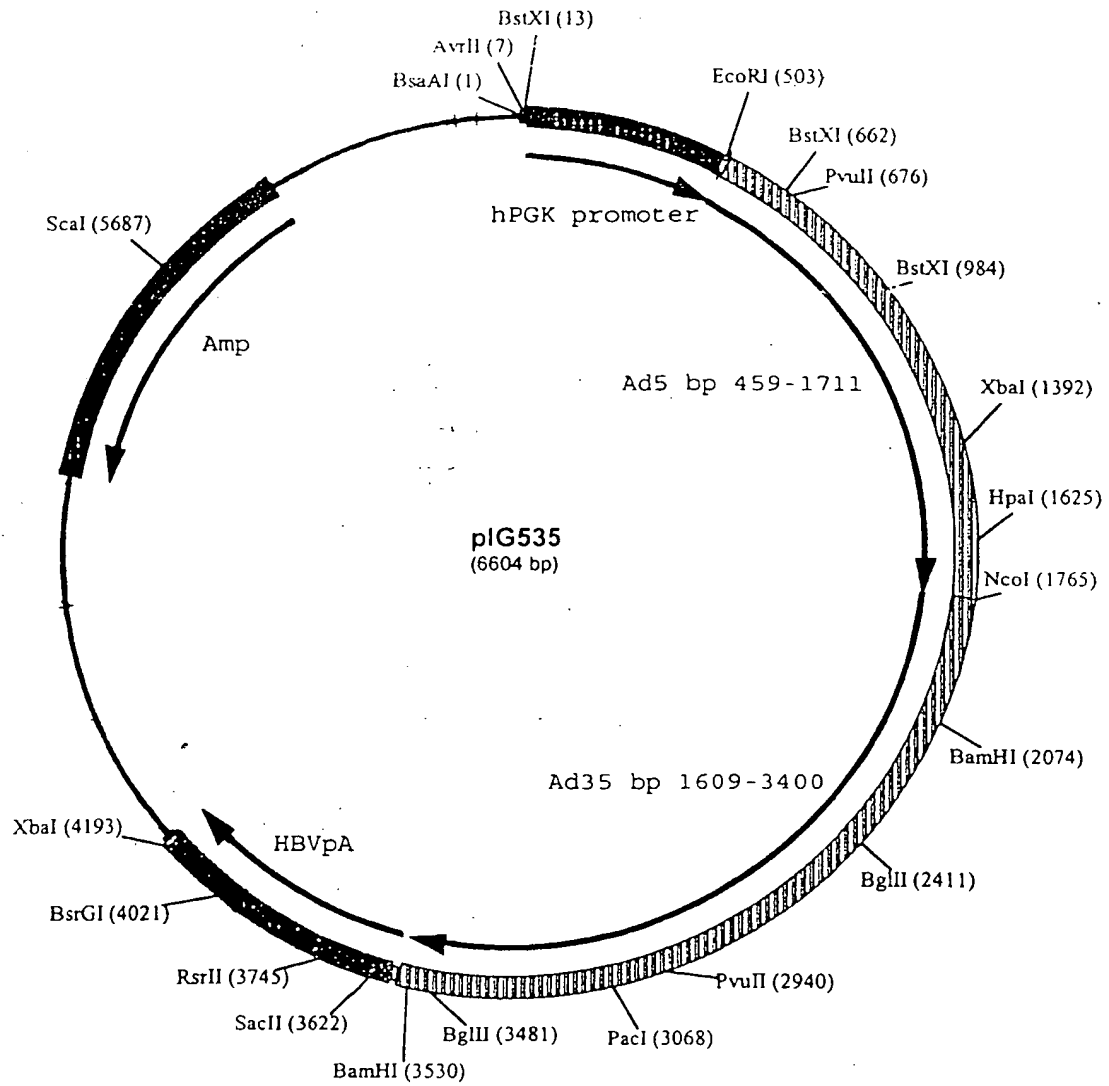


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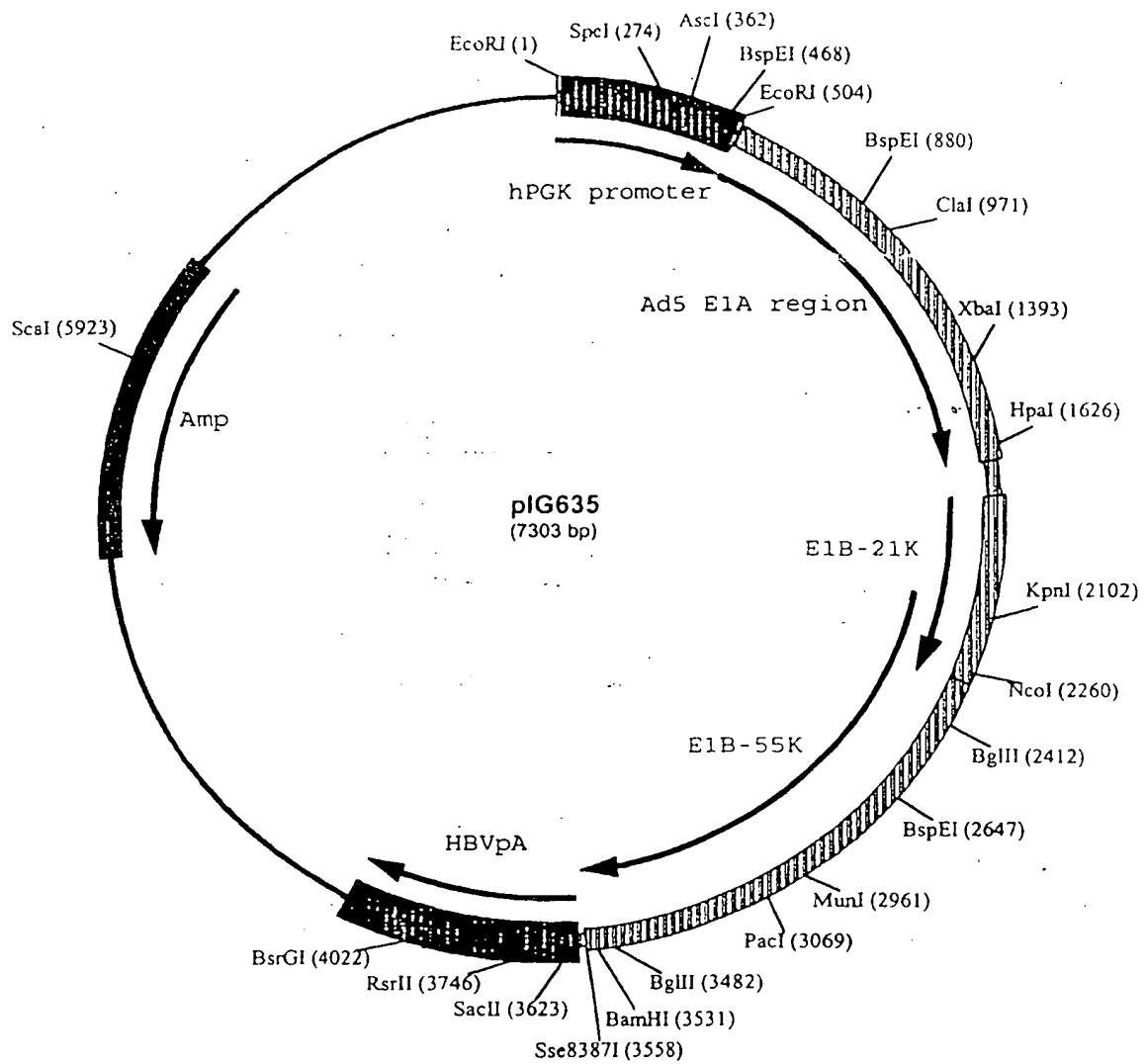


Figure 27

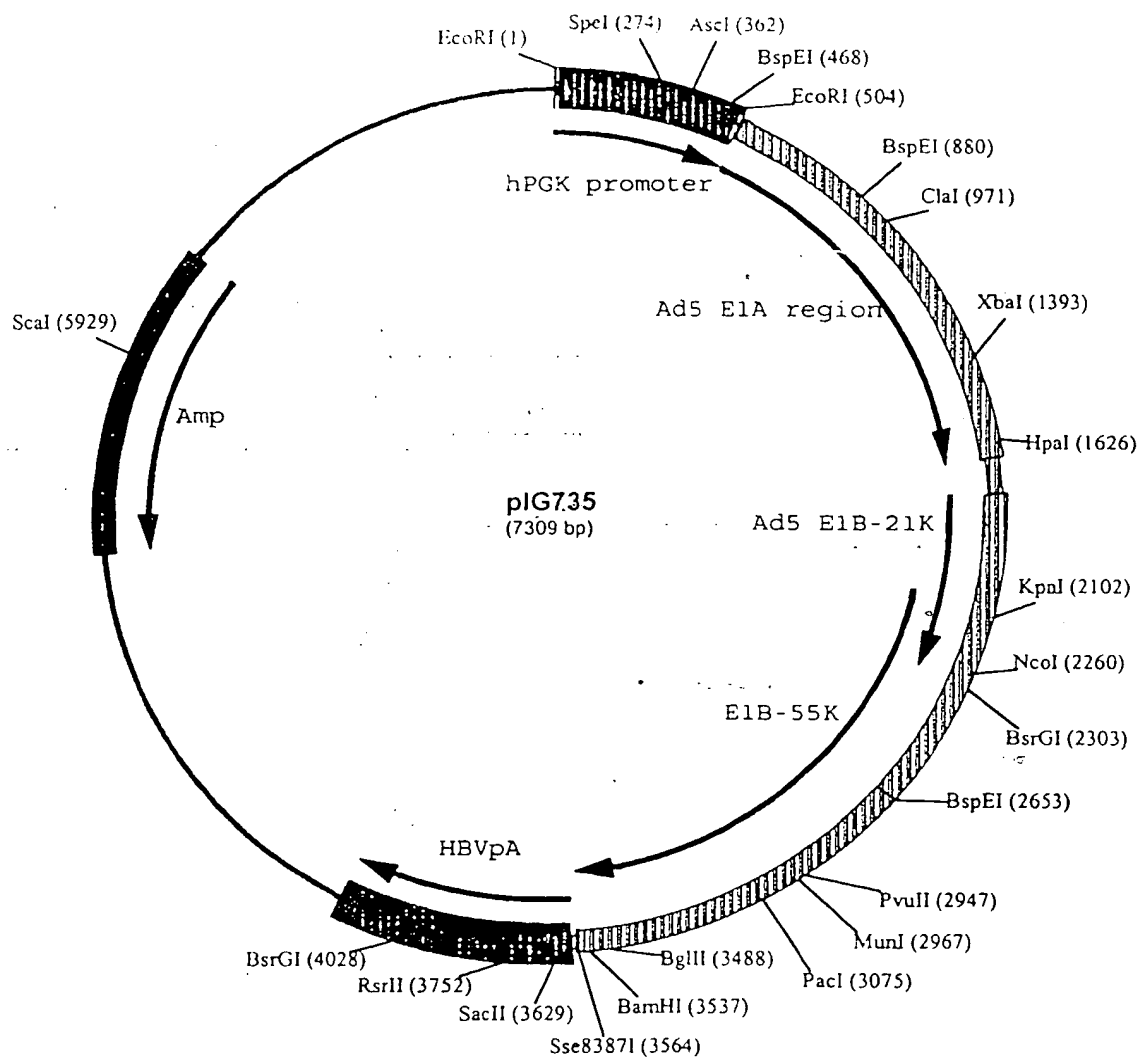


Figure 28

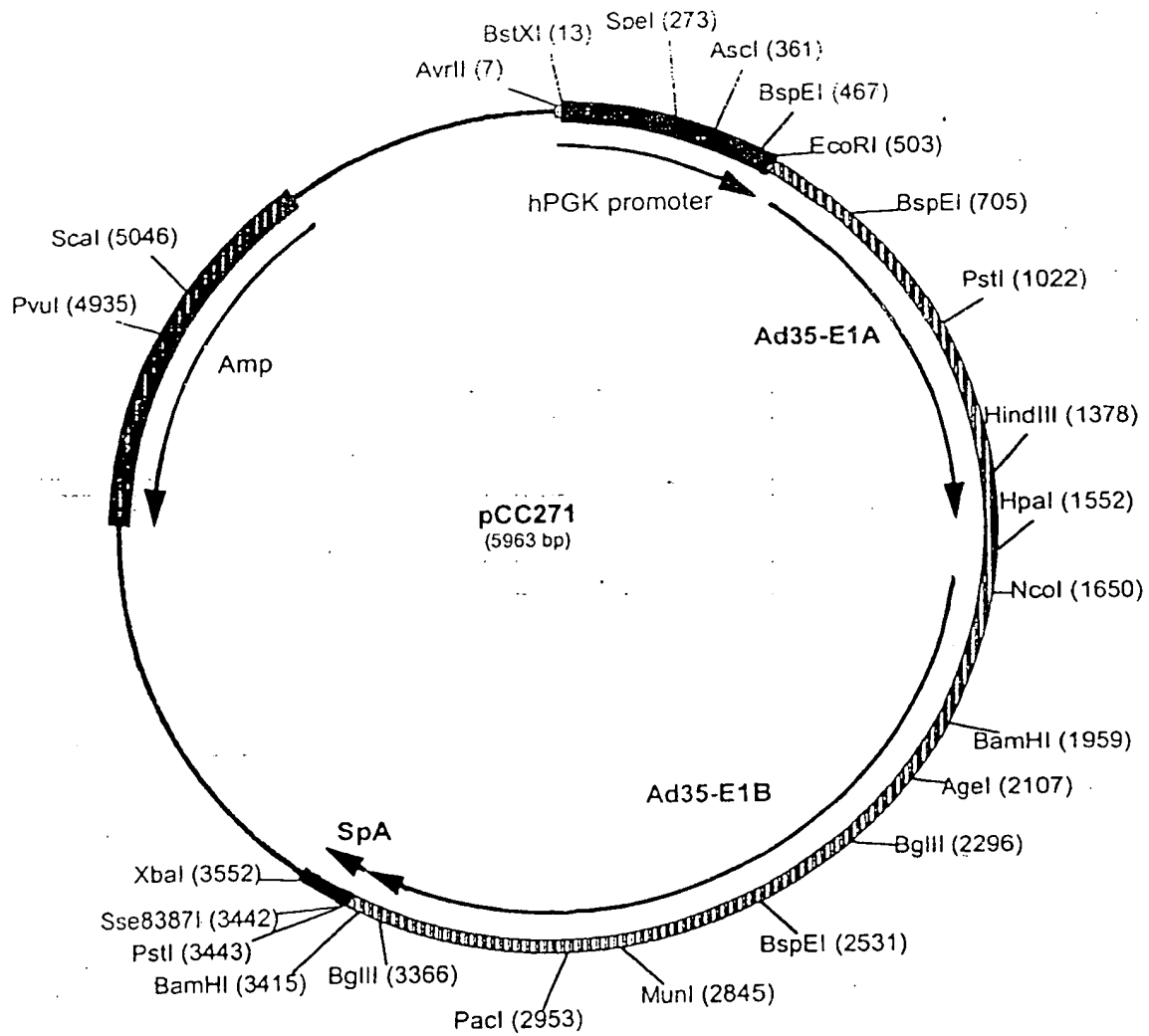


Figure 29

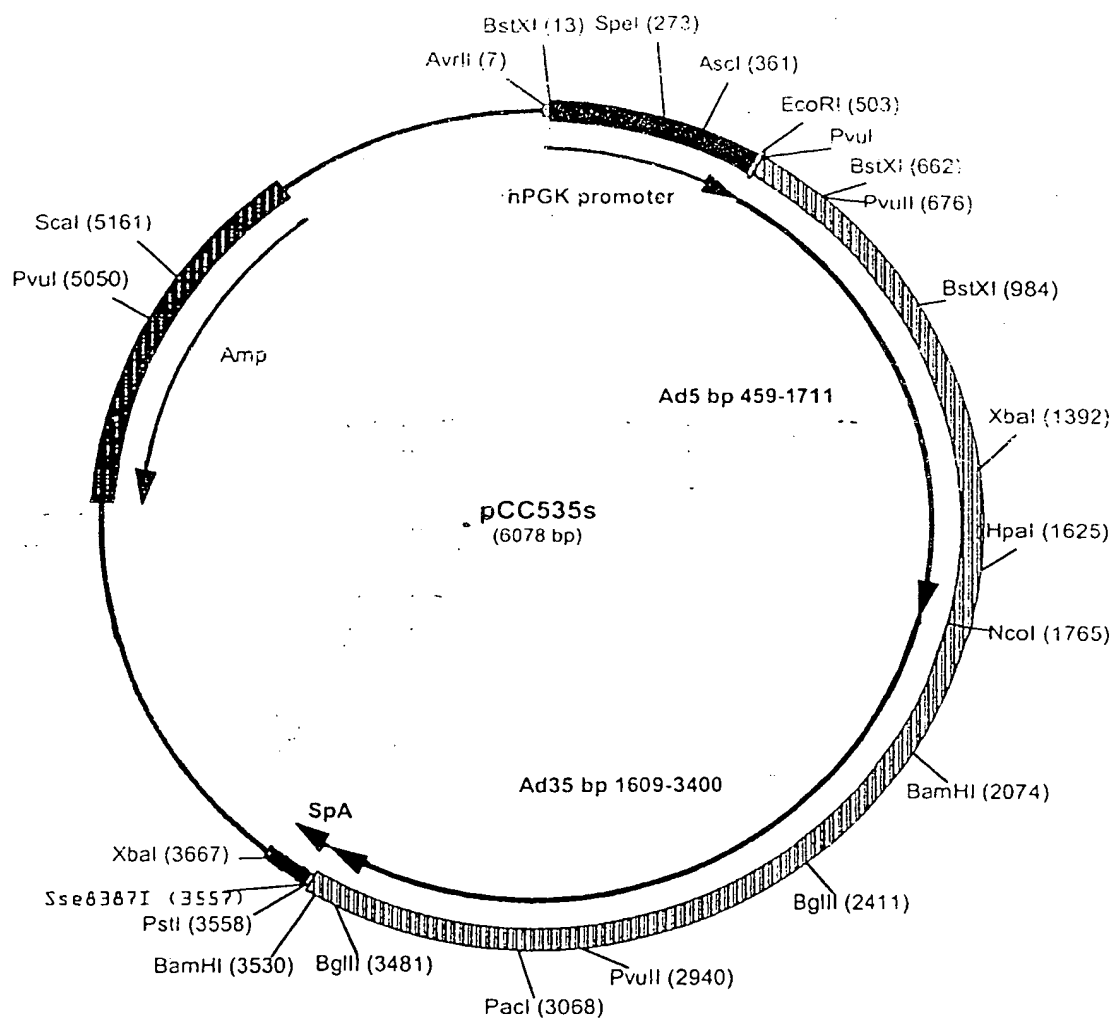


Figure 30

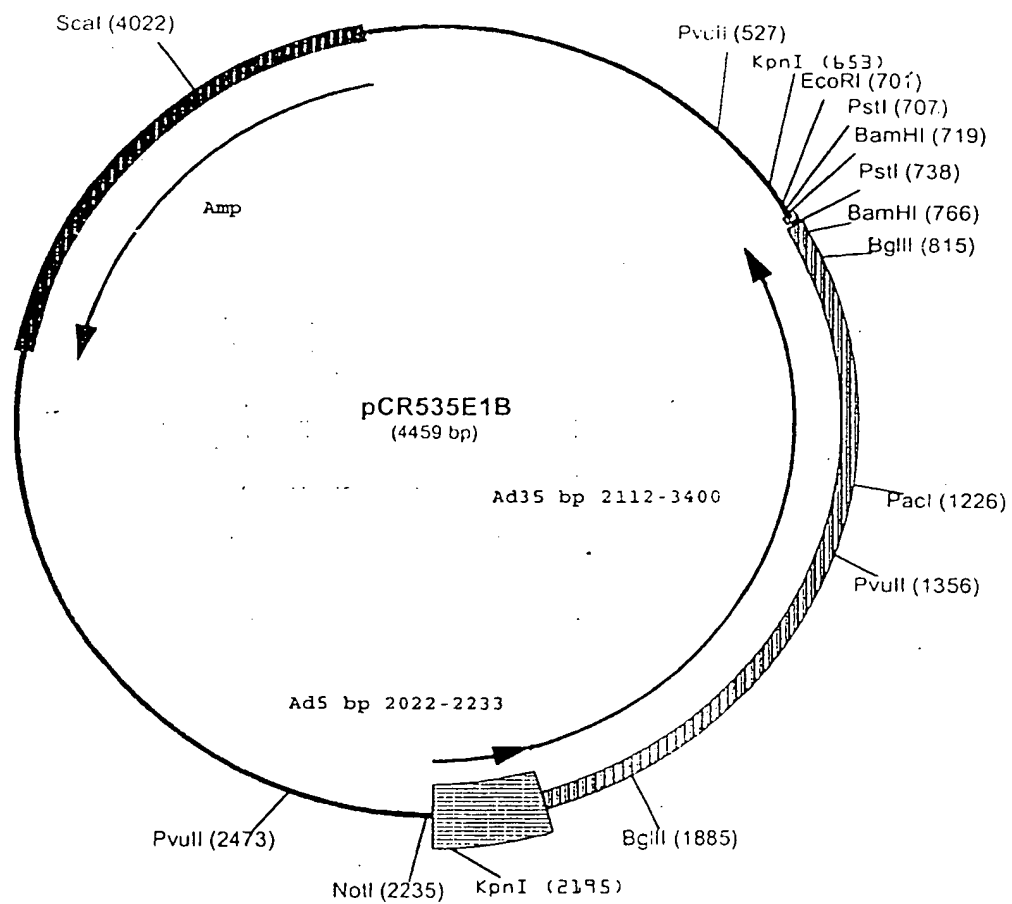


Figure 31

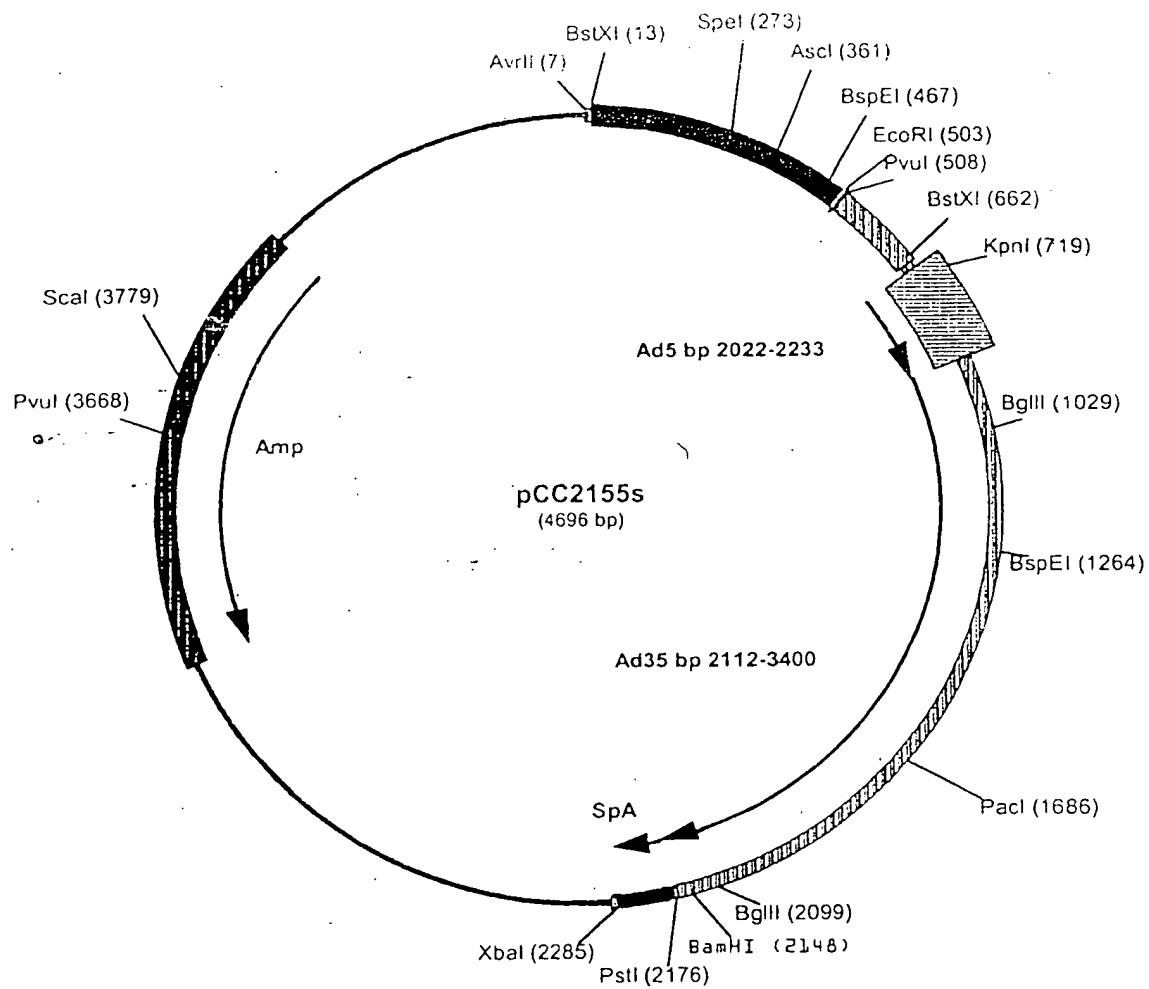


Figure 32

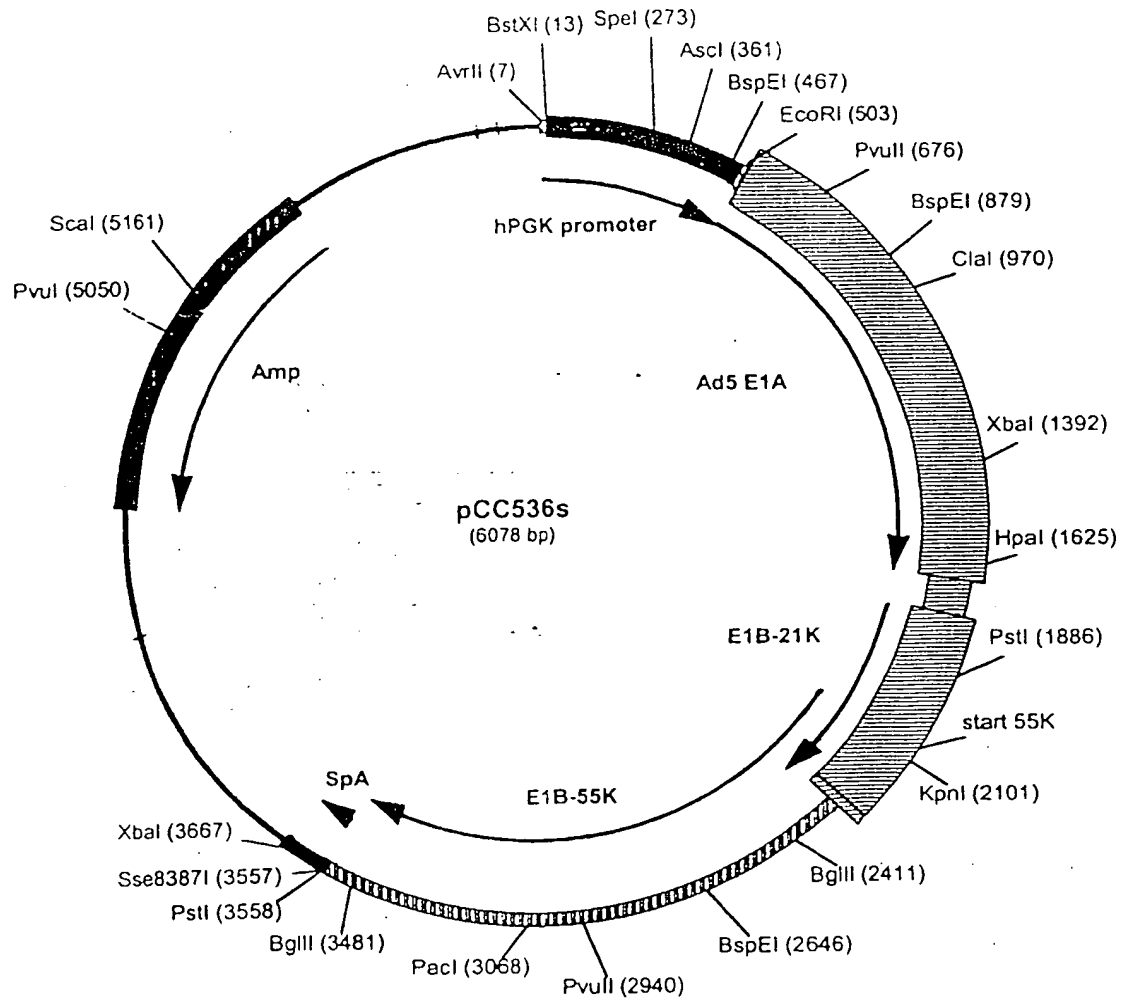


Figure 33

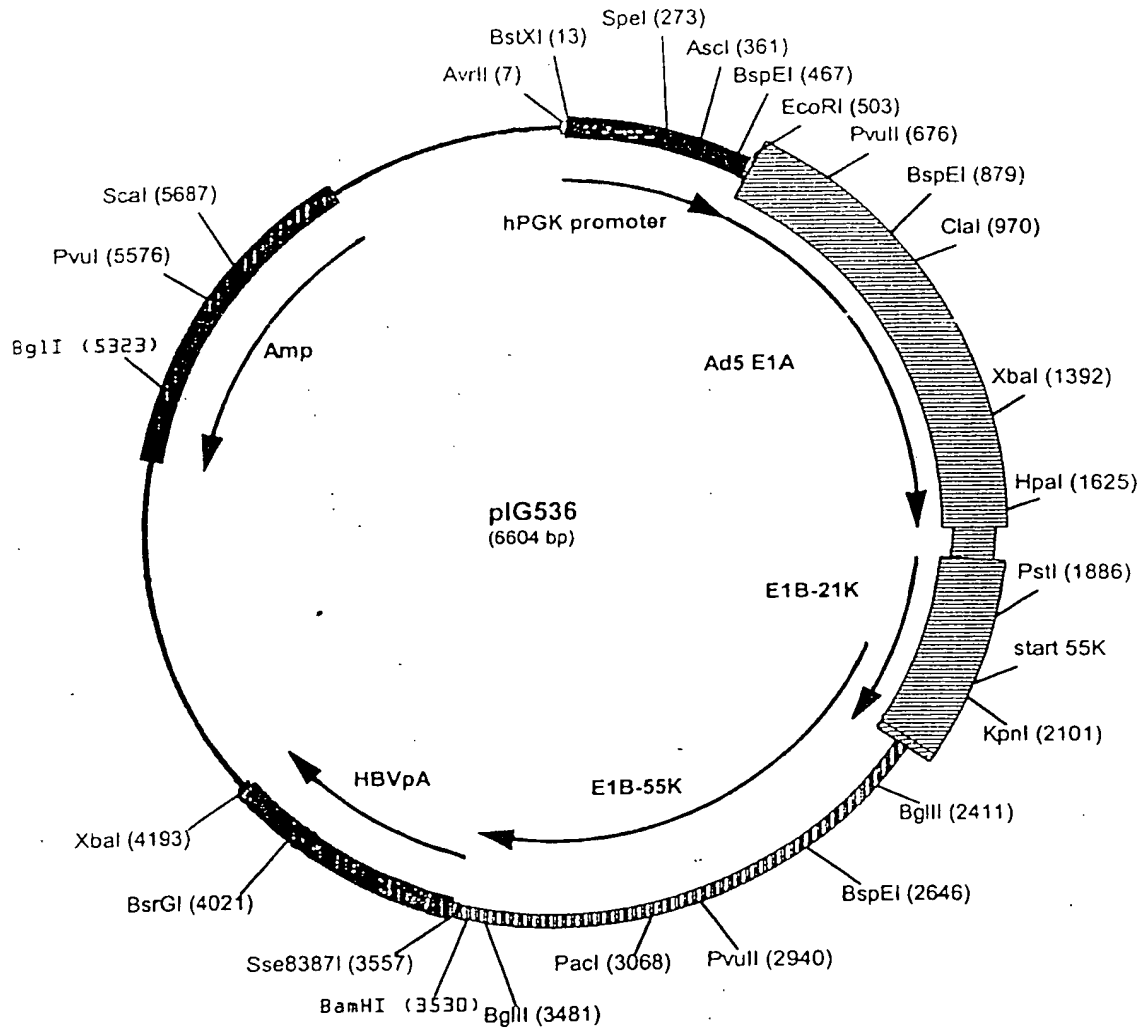


Figure 34

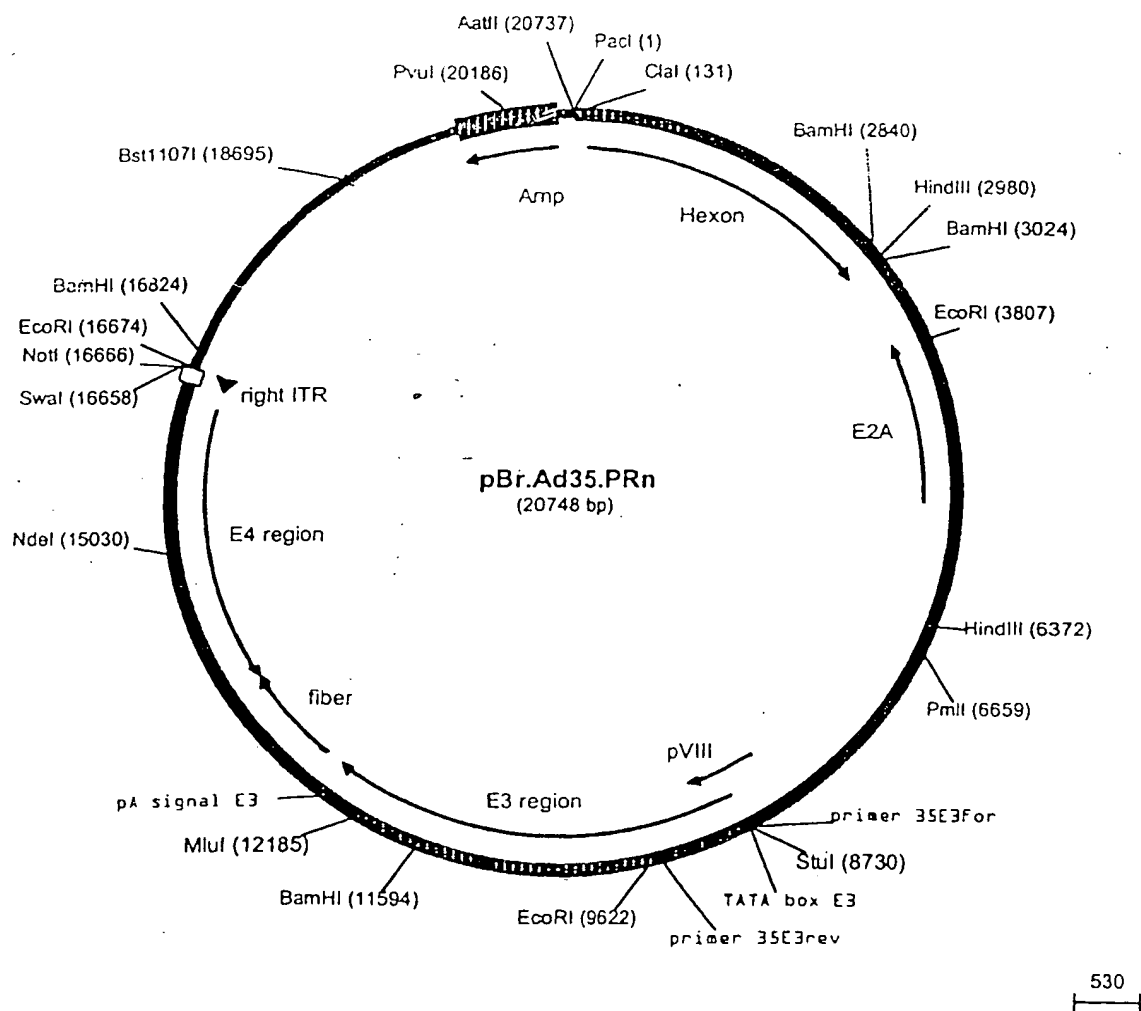


Figure 35

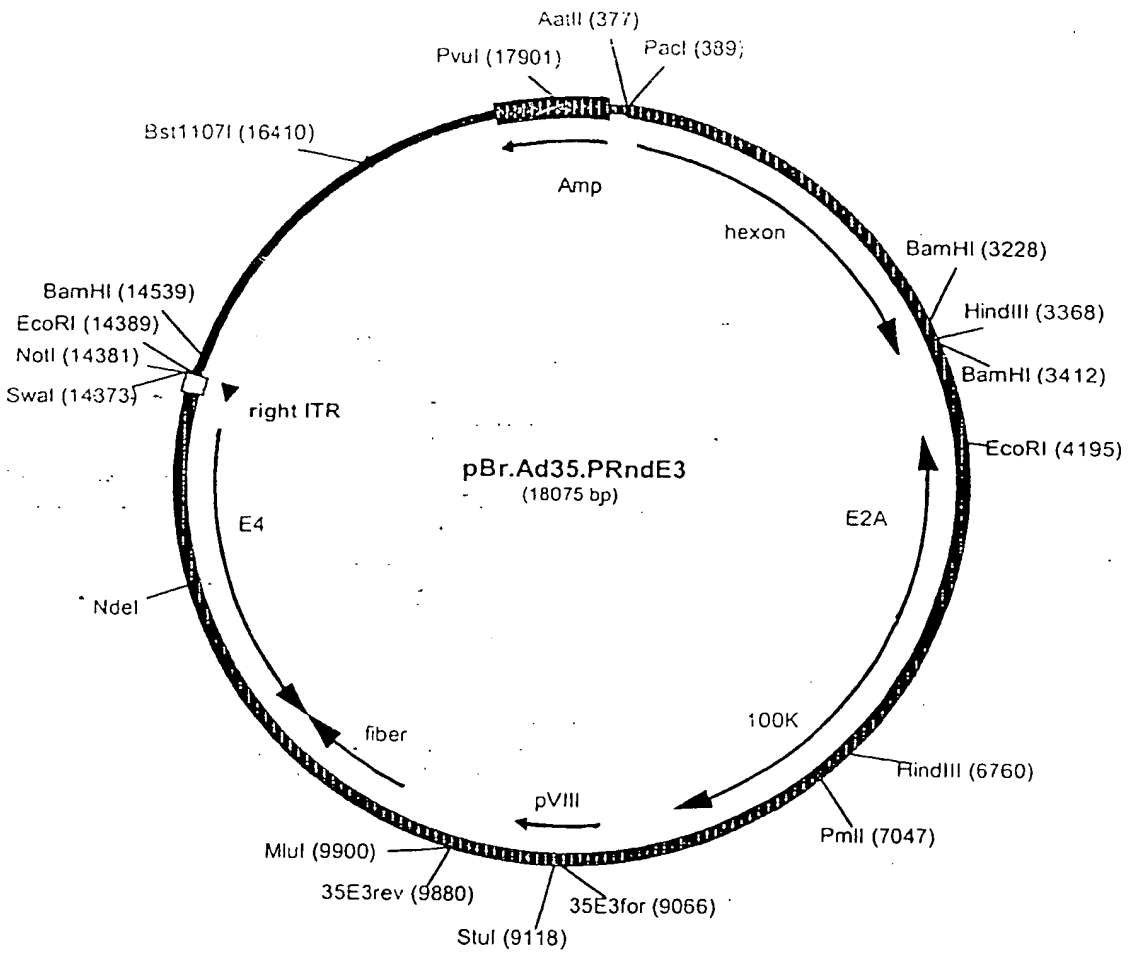
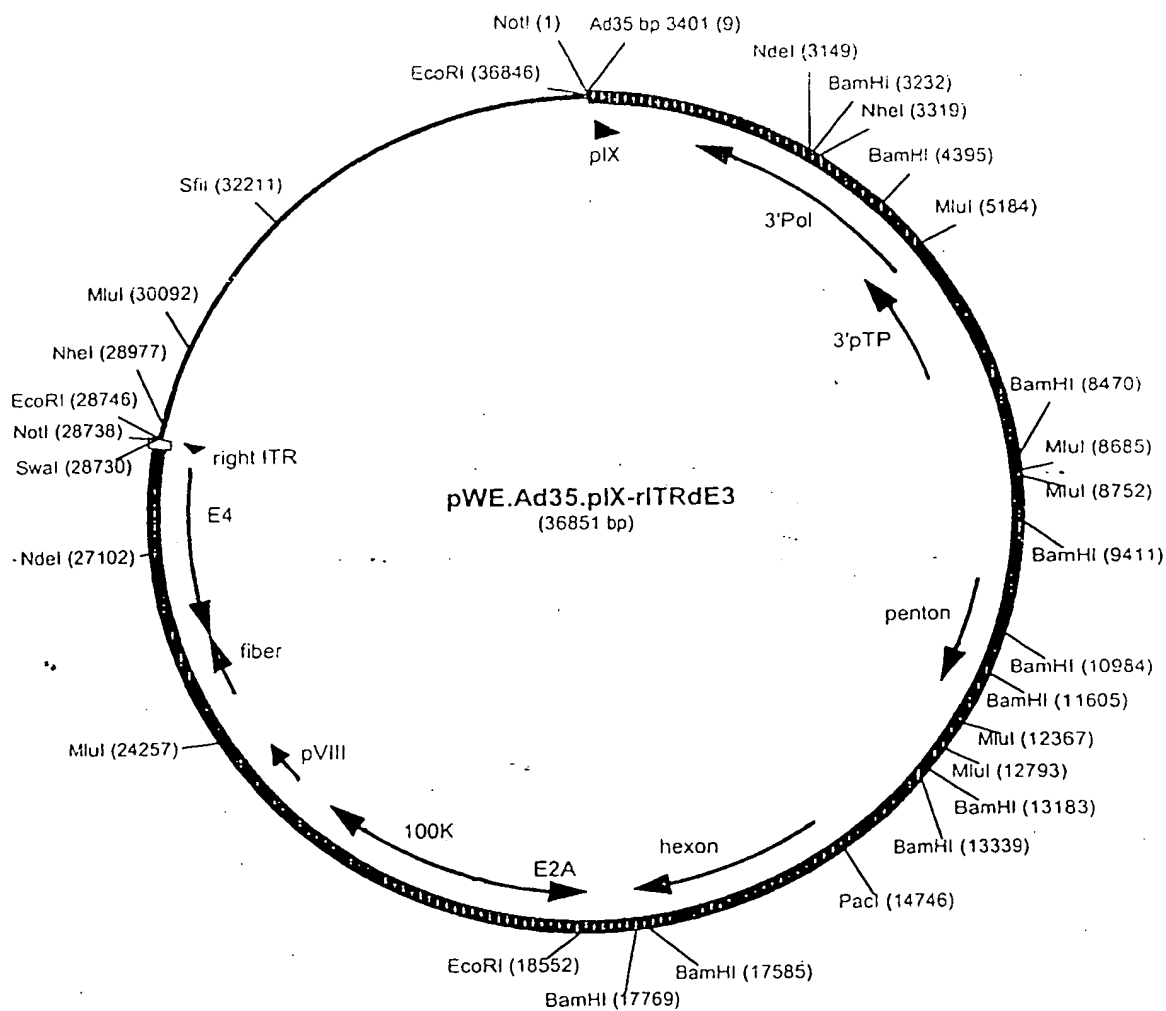


Figure 36



Inventor: Vogels et al.
Serial No.: 10/002,750
Docket No.: 2183-5148US

Figure 37 B: Alignment of E1B-55K sequences from pCC536s, wtAd35 and wtAd5

1 MERNPSERGVPA GFSGHASVESGCGTQESPATVVFRRPPGDNITDGGAAAAGGSQAAAAAG PCC536s.55K PRO
 1 MDPADSFOQ G I R F G F H S H S I V E N M E G S O D E D N L R L L A S A F G C S G N P E A S T G H A S G S G G G A D35 E1B-55K pro
 1 MERNPSERGVPA GFSGHASVESGCGTQESPATVVFRRPPGDNITDGGAAAAGGSQAAAAAG A45 E1B-55K pro
 61 AEPMEPESRPPGPPSS I GGGGVADLSPELQORVLTGSTSTGDRGVKKRERASS I GTDARSEL A PCC536s.55K PRO
 61 TARGOQ P E S R P P G P S S I GGGGVADLSPELQORVLTGSTSTGDRGVKKRERASS I GTDARSEL A A45 E1B-55K pro
 61 AEPMEPESRPPGPPSGMNVVQV A E L Y I P E L R I I L T I T E D G Q G L K G V K R E R G A C E A T E E A R N I L A A45 E1B-55K pro
 119 LSLMSRRRPETI WWHEVQKEGRDEVSVLQEKYSLEQVKT CWLEPEDD WAVA I K N Y A K I A L PCC536s.55K PRO
 119 LSLMSRRRPETI WWHEVQKEGRDEVSVLQEKYSLEQVKT CWLEPEDD WAVA I K N Y A K I A L A45 E1B-55K pro
 121 FSLMTTRH R P E C I T F Q O I K D N C A N E L D L A Q I K Y S I L E Q L T Y W L Q P G D D F E E A I R V Y A K V A L A45 E1B-55K pro
 179 RPKQYKISRRRI N I R N A C Y I S G N G A E V V I D T Q D K T V I R C C M M D M W P G V V G M E A V T F V N V K PCC536s.55K PRO
 179 RPKQYKISRRRI N I R N A C Y I S G N G A E V V I D T Q D K T V I R C C M M D M W P G V V G M E A V T F V N V K A45 E1B-55K pro
 181 RPDCKX Y K I S K L V N I R N C C Y I S G N G A E V E I D T E D R V A F R C S I M I N M W P G V L G M D G V V I M N V R A45 E1B-55K pro
 239 FRGDGYNGI VFMANT K L I L H G C S F F G F N N T C V D A W G Q V S V R G C S F Y A C W I A T A G R T K S Q L PCC536s.55K PRO
 239 FRGDGYNGI VFMANT K L I L H G C S F F G F N N T C V D A W G Q V S V R G C S F Y A C W I A T A G R T K S Q L A45 E1B-55K pro
 241 FTGPNFSGT V F L A N T N L I L H G V S F Y G F N N T C V E A W T D V R V R G C A F Y C C W K G V V C R P K S R A A45 E1B-55K pro
 269 SLKKCIFORCNLGI L N E G E A R V R H C A S T D T G C F I L I K G N A S V K H N M I C G A S D E R P Y Q M L T PCC536s.55K PRO
 299 SLKKCIFORCNLGI L N E G E A R V R H C A S T D T G C F I L I K G N A S V K H N M I C G A S D E R P Y Q M L T A45 E1B-55K pro
 301 S I K K C L F E R C T L G I L S E G N S R V R H N V A S D C G C F M L V K S V A V I K H N M V C G N C E D R A S Q M L T A45 E1B-55K pro
 359 CAGGHCNMLATVHI VSHQRKKWPVFDHNVLTKCTMHAGRRRGFMFMPYQCNCNMNHVKVLLLEP PCC536s.55K PRO
 359 CAGGHCNMLATVHI VSHQRKKWPVFDHNVLTKCTMHAGRRRGFMFMPYQCNCNMNHVKVLLLEP A45 E1B-55K pro
 361 CSDGNCHL L K T I H V A S H S R K K A W P V F E H N I L T R I C S L H L G N R R G V F L P Y O C N L S H T K T L L E P A45 E1B-55K pro
 419 OAFSRMSLTGIFDMNTQI WKI L R Y O D T R S R V R A C E C G G K H A R F Q P V C V D V T E D L R P D H L V PCC536s.55K PRO
 419 OAFSRMSLTGIFDMNTQI WKI L R Y O D T R S R V R A C E C G G K H A R F Q P V C V D V T E D L R P D H L V A45 E1B-55K pro
 421 E S M S K V N L N G V F O M T M K I W K V L R Y D E L T R T R C R P C E C G G K H I R N Q P V M L D V T E E L R P D H L V A45 E1B-55K pro
 479 IARTGA EFGSSGEETD 479 PCC536s.55K PRO
 479 IARTGA EFGSSGEETD A45 E1B-55K pro
 481 L A C I T R I A E F G S S D E D T D 481 A45 E1B-55K pro

Decoration 'Decoration #1': Box residues that differ from pCC536s.55K.PRO.

Figure 37 A: Alignment of E1B-21K sequences from pCC536s, wtAd35 and wtAd5

1 MEAWECLEDFSAVRNLLLEQSSNSTSWFWRFLWGSQAQKLVCRIRKEDYKWE pCC536s.21K.pro
1 MEAWECLEDFSAVRNLLLEQSSNSTSWFWRFLWGSQAQKLVCRIRKEDYKWE Ad5.E1B-21K.pro
1 MEVWAILEDLRKTRQLLESASDGVSGPWRFWFASELARVFRIKQDYKQE Ad35.E1B-21K.pro
51 FEELKSCGELFDLSNLGHAALFQEKVIXITLDFSTPGRAAAAVAFLSPIK pCC536s.21K.pro
51 FEELKSCGELFDLSNLGHAALFQEKVIXITLDFSTPGRAAAAVAFLSPIK Ad5.E1B-21K.pro
51 FEELKSCGELFDLSNLGHAALFQEKVIXITLDFSTPGRAAAAVAFLSPIK Ad35.E1B-21K.pro
101 DKWSEETHLSGGYLLDPLAMHLWRA-VVRRHKNRLLLSVVRPAIIPTEEQ pCC536s.21K.pro
101 DKWSEETHLSGGYLLDPLAMHLWRA-VVRRHKNRLLLSVVRPAIIPTEEQ Ad5.E1B-21K.pro
101 DKWIPQTHPSRGYVLDPLATLWRTWVVRKMRITLGYWVPVQLGVAGILR Ad35.E1B-21K.pro
150 QQQEEARRRRQEQSPWNPRAGLDPPVEEA pCC536s.21K.pro
150 QQQEEARRRRQEQSPWNPRAGLDPPVEEA Ad5.E1B-21K.pro
151 HPPVMPAVLEEEQVED-NPRAGLDPPVEEA Ad35.E1B-21K.pro

Decoration #1: Box residues that differ from the Consensus.